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PROVISIONAL APPLICATION FOR PATENT COVER SHEET
This is a request for filing a PROVISIONAL APPLICATION FOR PATENT under 37 CFR §1.53(c).

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Attorney's Docket No.: 15138-003P01

# PROVISIONAL APPLICATION FOR PATENT

# under

# 37 CFR §1.53(c)

TITLE:

METHODS AND COMPOSITIONS FOR DISEASE

**DIAGNOSIS** 

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# METHODS AND COMPOSITIONS FOR DISEASE DIAGNOSIS

## FIELD OF THE INVENTION.

The invention relates to methods and compositions for risk assessment,

identification, diagnosis, prognosis, and/or monitoring of disease, and for early therapeutic intervention.

# BACKGROUND OF THE INVENTION

It is axiomatic that early diagnosis and concomitant early therapeutic intervention is the key to successful treatment and/or management of most human disorders. However, many disorders cannot be diagnosed until the pathological process is already advanced. For example, many solid tumors are usually not clinically detectable before they can be palpated or visualized by tissue imaging techniques (i.e., when they are at least 0.5 cm in size), at which time neoplasia may have been present for years. Similarly, the diagnostic criterion for diabetes mellitus (increased fasting plasma glucose levels or hyperglycemia) identifies the disorder when glucose intolerance (the underlying cause of hyperglycemia) is already present. In another example, rheumatoid arthritis (RA) is diagnosed by the presence of joint stiffness and soreness and the presence of positive rheumatoid factor, all factors that indicate RA is already present and may be advanced.

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#### Diagnostic Disease Markers

In cancer, progression from preneoplasia to malignancy is accompanied by the accumulation of genetic changes in the neoplastic cells that lead to histopathological modifications. In some circumstances, when such a genetic change corresponds to an increase in a protein made by the tumor cells, such a protein can be detected in the tumor or in body fluids (if secreted from the tumor), and used as a biological tumor marker. Most tumors have been associated with one or more such tumor markers. Such markers have been evaluated as potential tools to diagnose cancer, determine prognosis, and/or monitor cancer progression. However, many tumor markers are detectable only after neoplasia has already progressed to the stage of formation of a tumor. In some cases, a tumor marker may not be detectable until a tumor is already malignant. Thus, many of

the most widely used tumor markers are used primarily to monitor disease progression or response to treatment rather than for early diagnosis.

In rheumatoid arthritis, anti-cyclic citrullinated peptide (anti-CCP) antibodies, anti-keratin antibodies (AKA) and IgM rheumatoid factors have been suggested as markers for rheumatoid arthritis (Bas et al., Rheumatology (Oxford), 2002, 41(7):809-14). However, the value of such markers remains inconclusive (Scott, Rheumatology (Oxford), 2000, 39(Supp) 1:24-9). Similarly, while several protein and gene markers have been found to correlate with the presence of active diabetes, the use of markers as diagnostic or predictive has not been proven valuable at this time for either type I or type 2 diabetes (see the National Academy of Clinical Biochemistry (NACB) Laboratory Medicine Practice Guidelines: Guidelines and Recommendations for Laboratory Analysis in the Diagnosis and Management of Diabetes Mellitus, 2002, available online at the NACB web site).

# 15 Genomics and Proteomics Tools for Disease Diagnosis

The development of high throughput screening approaches such as functional genomics and proteomics has provided a new biological platform to search for molecules associated with different disorders. Gene-expression profiles based on microarray analysis have been of some use to predict survival of patients with lung carcinoma (Beer et al., 2002, Nat. Med., 8(8):816-24). A similar approach identified a group of genes that were said to be useful to predict the clinical outcome of diffuse large B-cell lymphoma following combination chemotherapy (Shipp et al., 2002, Nat. Med., 8(1):68-74). In addition, comparison of the proteomic profile of patients with ovary or prostate cancer compared to non cancerous volunteers was said to have provided a set of serum proteins that might be useful for early cancer detection (Petricoin et al., 2002, Lancet, 2002, 359(9306):572-7; Petricoin et al., 2002, J. Natl. Cancer Inst., 94(20):1576-8).

At present, most functional genomics studies in cancer have used cancer samples obtained from patients to generated cancer-associated gene expression profiles (either by a genomics or a proteomics approach).

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A need remains for methods to detect and diagnose disease. Particularly needed are predictive methods and markers for early stage or very early stage disease detection and risk assessment.

#### **SUMMARY OF THE INVENTION**

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The methods described herein are based, at least in part, on the discovery that the central nervous system (CNS) exhibits specific changes in gene expression (e.g., changes in patterns of gene expression) in response to the presence of a peripheral (non-CNS) disease or disorder (e.g., a hyperproliferative disorder such as a non-CNS tumor or cancer, an immunological disorder, an inflammatory disorder, a metabolic disorder, or a pathogenic infection). While not bound by any theory, the inventors believe that specific changes in gene expression in the CNS, e.g., in the brain, occur in response to the presence of peripheral disease at an early stage in the development of the disease, e.g., before the disorder is clinically detectable and/or before the subject is symptomatic. Thus, peripheral disorders can be diagnosed at an early stage and targeted for early therapeutic intervention by analyzing changes or patterns in gene expression in the CNS.

Accordingly, in one aspect, the invention features a method of diagnosing a non-CNS disorder in a subject, such as a human. The non-CNS disorder can be, e.g., a hyperproliferative disorder, e.g., a non-CNS tumor or cancer; an immunological disorder, e.g., rheumatoid arthritis; an inflammatory or allergic disorder, e.g., asthma; a metabolic disorder, e.g., diabetes or obesity; or a pathogenic infection, e.g., a viral infection. The method includes detecting expression of a gene in a CNS sample of the subject, e.g., a brain tissue or cell (such as a tissue or cell of the hypothalamus, the cerebellum, the midbrain, the hippocampus, the prefrontal cortex or the striatum) or a sample of cerebrospinal fluid (CSF). The method optionally includes a step of obtaining the CNS sample. A change in gene expression compared to a reference value, e.g., a control or basal value, is correlated with the presence of a non-CNS disorder. The method is not limiting in that it can be used to detect the risk or presence of any non-CNS disorder. In one embodiment, the non-CNS disorder is not lymphoma.

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The subject can be a human. In one embodiment, the human is not symptomatic for the disorder to be diagnosed. In another embodiment, the disorder is not clinically detectable, e.g., it is not detectable by a routine general clinical exam.

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Detecting expression of a gene in a CNS sample can include detecting or determining a value for one or more of: the level of mRNA, rate of transcription, amount of a gene product, and activity of a gene product. In some embodiments, expression of a single gene in the CNS may be detected, where a change in gene expression in that gene is associated with the presence of a non-CNS disorder. In other embodiments, expression of a plurality of genes (e.g., a panel or cluster of genes) may be evaluated, where a specific profile of gene expression of the plurality of genes is associated with the presence of a particular non-CNS disorder.

The method can include correlating the result of the detecting step to the presence or absence of a non-CNS disorder. "Correlating" means identifying the probability, based on the result of the detecting step, that the subject has or does not have a non-CNS disorder. Correlating can include generating a dataset from, or providing a record of, the detecting step, e.g., a printed or computer readable record such as a laboratory record or dataset. The record can include other information, such as a specific subject identifier, a sample identifier for the CNS sample, a date, the identity of the operator of the method, and/or other information. The record can be used to provide or store information about the subject. For example, the record can be used to provide information (e.g., to the subject, a health care provider, the government, or insurance company). The record or information derived from the record can be used, e.g., to identify the subject as suitable or unsuitable for a particular therapy or a particular clinical trial group.

In the methods described herein, gene expression of a CNS gene can be detected by any technique available to the skilled artisan, e.g., genomics or proteomics microarray analysis of a CNS biological sample, such as brain tissue or CSF; or brain imaging techniques that detect changes in gene expression. In one embodiment, the method involves detecting a CNS gene product released or secreted into the CSF. In such embodiments, an agent (such as an antibody, e.g., a labeled antibody) for detecting the gene product can be immobilized on a solid phase, e.g., in a dipstick format.

The gene or genes to be evaluated will depend on the specific gene or profile of gene expression associated with a particular disorder. For example, exemplary genes (or profiles or clusters of genes) that are regulated in response to the presence of cancer cells (or particular types of cancer cells) are shown in FIGS. 2-26, *infra*. Such genes are also referred to herein as CNS "marker genes" or "disease surveillance genes" for non-CNS disorders. The exemplary CNS marker genes are not limiting, as the methods described herein can include the detection of any other genes or gene products determined to exhibit a change in expression associated with the presence of a peripheral disorder. CNS marker genes can include, *inter alia*, genes encoding hormones, growth factors, immune system components, and cytokines.

In another aspect, the invention features a method of determining whether a subject (e.g., a human) has, or is at risk for developing, a peripheral (non-CNS) disorder. The method involves providing or obtaining a test gene expression profile for two or more CNS genes in the subject; and comparing the test gene expression profile with a reference gene expression profile (e.g., a reference gene expression profile described herein), wherein the reference gene expression profile is associated with the presence of a particular non-CNS disorder. Non-limiting examples of reference gene expression profiles (e.g., associated with colon, breast or lung carcinoma), are disclosed herein. In one embodiment, the method includes generating a record of the result (e.g., a laboratory record or dataset) of the comparing step; and, optionally, transmitting the record (e.g., by print or computer readable material) to the subject, the subject's health care provider or another party. As with other methods described herein, various techniques can be used to provide a gene expression profile and various types of disorder scan be detected.

In another aspect, the invention features a method of treating a subject, e.g., a human, by diagnosing a peripheral (non-CNS) disorder in a subject, e.g., using a method described herein; and administering to the subject a therapeutic agent for the treatment of the disorder, e.g., a chemotherapeutic agent. Because the detection/diagnostic methods described herein can indicate the presence of peripheral disease at an early stage in the pathogenic process (e.g., before a disorder is symptomatic or clinically diagnosable), the methods allow for early intervention to control the disorder, e.g., implementing lifestyle changes to stop or slow further progress of the disease, or by administering a therapeutic

agent to slow or control the progression of the disease. Such agents can advantageously be used at lower dosages than are typically used after a disease is sufficiently advanced to be clinically diagnosed.

In another aspect, the invention features a method of identifying a diagnostic marker gene for a peripheral (non-CNS) disorder in a subject. The method involves: inducing a non-CNS disorder in a test experimental animal, e.g., a rodent tumor model; comparing expression of a gene in a CNS tissue or cell in the test experimental animal to expression of the gene in a control experimental animal; and selecting as a diagnostic marker a gene (or human homolog of the gene) that is differentially expressed in the test experimental animal compared to the control experimental animal.

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The methods described herein are useful, *inter alia*, for risk assessment for a variety of disorders, for early detection and diagnosis of disease, for monitoring of progression of disease, for monitoring efficacy of treatment for a disease, and/or evaluation of clinical status.

As used herein a "disorder" or "disease" is an alteration in the state of the body or of some of its cells, tissues, or organs, that threatens health. The two terms are meant to encompass all stages of an illness, including the very early stages of an illness (e.g., early alterations in the body that may not be detectable by the subject or a health care provider, but nonetheless set in motion a disease process). For example, the terms "disorder" and "disease" encompass the state of neoplasia, before a neoplasm or tumor is formed; early immunological reactions to an antigen, e.g., in the development of rheumatoid arthritis or asthma, before inflammation or allergy are symptomatic; and early changes in energy metabolism that promote weight gain, before weight gain is produced.

As used herein, "neoplasia" is an unregulated and progressive proliferation of cells under conditions that would not elicit, or would cause cessation of, proliferation of normal cells. Neoplasia can result in the formation of a "neoplasm," a new and abnormal growth of tissue. If the abnormally proliferating cells form a mass, a neoplasm is generally referred to as a "tumor." A neoplasm may be benign or malignant (cancerous).

As used herein, the term "matches" or "matching" when referring to a test gene expression profile and a reference gene expression profile, means that the profiles are sufficiently similar to each other to have an analogous cause or effect. Two profiles are

matching if they are at least 70% identical in reference to the number of genes having similar expression patterns in each profile, or the level of expression of the genes in each profile. In some embodiments, two profiles can be at least 80%, 85%, 90%, 95%, 98%, or more, identical.

A "subject" is a human or animal that is tested for the presence of a possible disorder. The animal can be a mammal, e.g., a domesticated animal such as a dog, cat, horse, pig, cow or goat; an experimental animal such as an experimental rodent (e.g., a mouse, rat, guinea pig, or hamster); a rabbit; or an experimental primate, e.g., a chimpanzee or monkey.

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Although methods and materials similar or equivalent to those described herein can be used in the practice or testing of the present invention, suitable methods and materials are described below. All publications, patent applications, patents, and other references mentioned herein are incorporated by reference in their entirety. In case of conflict, the present specification, including definitions, will control. In addition, the materials, methods, and examples are illustrative only and not intended to be limiting.

Other features and advantages of the invention will be apparent from the following detailed description, the drawings, and the claims.

#### BRIEF DESCRIPTION OF THE DRAWINGS

FIG. 1 is a schematic diagram of a slide used in a microarray gene expression assay.

FIG. 2-(1-7) is a table showing the results of cluster analysis I for colon cancer in midbrain. This cluster analysis identified differentially expressed genes (p < 0.05) up- or down-regulated at one of two experimental time points after injection of cancer cells into the relevant animal model. Genes with similar expression pattern were clustered using hierarchical clustering techniques. The table lists 407 markers differentially expressed in the midbrain of mice injected with colon cancer cells at one of the two experimental time points studied for colon cancer (72 and 198 hours). Each listed marker gene belongs to one of 12 clusters, as indicated by the left hand column of the table. FIG. 2-8 is a set of cluster graphs that illustrates the relative differential expression of each cluster at each of

the time points tested. In each cluster diagram (in all the figures), the y-axis represents a value for the relative level of gene expression of the cluster compared to control (the midline at 0) and the x-axis represents time. In this case, 1.0 = 72 hours and 2.0 = 198 hours. Thus, expression values below the midline indicates the genes in that cluster are down-regulated at that time. Expression values above the midline indicates the genes in that cluster are down-regulated at that time.

FIG. 3-1 is a table showing the results of cluster analysis II for colon cancer in midbrain. Cluster analysis II identified differentially expressed genes (p < 0.05) up- or down-regulated at both experimental time points tested (72 and 198 hours) after injection of cancer cells into the relevant animal model. The table lists 41 markers (in 12 clusters) differentially expressed in the midbrain of mice injected with colon cancer cells at both 72 and 198 hours. FIG. 3-2 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control at each time point.

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FIG. 4-(1-15) is a table showing the results of cluster analysis I for breast cancer in midbrain. The table lists 698 markers (in 12 clusters) differentially expressed in the midbrain of mice injected with breast cancer cells at one of the three experimental time points studied (18, 72 and 198 hours). FIG. 4-16 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

FIG. 5-(1-6) is a table showing the results of cluster analysis II for breast cancer in midbrain. The table lists 299 markers (in 12 clusters) differentially expressed in the midbrain of mice injected with breast cancer cells at two or three of the experimental time points studied (18, 72 and 198 hours). FIG. 5-7 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

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FIG. 6-(1-14) is a table showing the results of cluster analysis I for lung cancer in midbrain. The table lists 797 markers (in 12 clusters) differentially expressed in the midbrain of mice injected with lung cancer cells at one of the three experimental time points studied. FIG. 6-15 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

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FIG. 7-(1-4) is a table showing the results of cluster analysis II for lung cancer in midbrain. The table lists 230 markers (in 12 clusters) expressed in the midbrain of mice injected with lung cancer cells at two or three of the three experimental time points studied. FIG. 7-5 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours) and 3.0 = 198 hours) (x-axis).

FIG. 8-(1-12) is a table showing the results of cluster analysis I for colon cancer in cortex. The table lists 688 markers (in 12 clusters) differentially expressed in the cortex of mice injected with colon cancer cells at one of two experimental time points studied. FIG. 8-13 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the two time points (1.0 =72 hours and 2.0 = 198 hours) (x-axis).

FIG. 9-(1-2) is a table showing the results of cluster analysis II for colon cancer in cortex. The table lists 58 markers (in 12 clusters) differentially expressed in the cortex of mice injected with colon cancer cells at two or three of the three experimental time points studied. FIG. 9-3 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

FIG. 10-(1-12) is a table showing the results of cluster analysis I for breast cancer in cortex. The table lists 744 markers (in 12 clusters) differentially expressed in the cortex of mice injected with breast cancer cells in one of the three experimental time

points studied. FIG 10-13 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

FIG. 11-(1-5) is a table showing the results of cluster analysis II for breast cancer in cortex. The table lists 272 markers (in 12 clusters) differentially expressed in the cortex of mice injected with breast cancer cells at two or three of the three experimental time points studied. FIG 11-6 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis). 10

FIG.12-(1-12) is a table showing the results of cluster analysis I for lung cancer in cortex. The table lists 828 markers (in 12 clusters) differentially expressed in the cortex of mice injected with lung cancer cells at one of the three experimental time points studied. FIG 12-13 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

FIG 13-(1-5) is a table showing the results of cluster analysis II for lung cancer in cortex. The table lists 311 markers (in 12 clusters) differentially expressed in the cortex of mice injected with lung cancer cells at two or three of the three experimental time points studied. FIG. 13-6 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

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FIG. 14-(1-7) is a table showing the results of cluster analysis I for colon cancer in striatum. The table lists 361 markers (in 12 clusters) differentially expressed in the striatum of mice injected with colon cancer cells at one of the two experimental time points studied. FIG. 14-8 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the two time points (1.0 = 72 hours and 2.0 = 198 hours) (x-axis).

FIG 15-1 is a table showing the results of cluster analysis II for colon cancer in striatum. The table lists 40 markers (in 12 clusters) differentially expressed in the striatum of mice injected with colon cancer cells in both experimental time points studied. FIG 15-2 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the two time points (1.0=72 hours and 2.0 = 198 hours) (x-axis).

FIG 16-(1-8) is a table showing the results of cluster analysis I for lung cancer in striatum. The table lists 483 markers (in 12 clusters) differentially expressed in the striatum of mice injected with lung cancer cells at one of the three experimental time points studied. FIG 16-9 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

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FIG 17-(1-4) is a table showing the results of cluster analysis II for lung cancer in striatum. The table lists 234 markers (in 12 clusters) differentially expressed in the striatum of mice injected with lung cancer cells at two or three of the three experimental time points studied. FIG 17-5 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

FIG. 18-(1-7) is a table showing the results of cluster analysis I for colon cancer in hypothalamus. The table lists 389 markers (in 12 clusters) differentially expressed in the hypothalamus of mice injected with colon cancer cells in one of the two experimental time points studied. FIG. 18-8 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the time points (1.0=72 hours and 2.0 = 198 hours) (x-axis).

FIG. 19-1 is a table showing the results of cluster analysis II for colon cancer in hypothalamus. The table lists 51 markers(in 12 clusters) differentially expressed in the

hypothalamus of mice injected with colon cancer cells in both experimental time points studied. FIG. 19-2 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

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FIG. 20-(1-20) is a table showing the results of cluster analysis I for breast cancer in hypothalamus. The table lists 1252 markers (in 12 clusters) differentially expressed in the hypothalamus of mice injected with breast cancer cells at one of the three experimental time points studied. FIG. 20-21 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

FIG. 21-(1-6) is a table showing the results of cluster analysis II for breast cancer in hypothalamus. The table lists 366 markers (in 12 clusters) differentially expressed in the hypothalamus of mice injected with breast cancer cells at two or three of the three experimental time points studied. FIG. 21-7 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

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FIG. 22-(1-20) is a table showing the results of cluster analysis I for lung cancer in hypothalamus. The table lists 1160 markers(in 12 clusters) differentially expressed in the hypothalamus of mice injected with lung cancer cells at one of the three experimental time points studied. FIG. 22-21 is a set of cluster graphs that illustrates the relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

FIG. 23-(1-6) is a table showing the results of cluster analysis II for lung cancer in hypothalamus. The table lists 306 markers (in 12 clusters) differentially expressed in the hypothalamus of mice injected with lung cancer cells at two or three of the three experimental time points studied. FIG. 23-7 is a set of cluster graphs that illustrates the

relative differential expression of each cluster compared to control (y-axis) at each of the three time points (1.0 = 18 hours, 2.0 = 72 hours and 3.0 = 198 hours) (x-axis).

FIG. 24(A)-(C) is a set of tables listing tumor-specific CNS markers differentially expressed, at any time tested, in three different cancer models: colon cancer, 24A; breast cancer, 24B; and lung cancer, 24C. Criteria for inclusion in this figure were (1) the marker corresponds to a secreted product; and (2) a p value below 0.05 for differential expression.

FIG. 25(A)-(E) is a set of tables listing genes identified as CNS markers that are also potential targets for therapeutic intervention for each of colon, breast and lung cancer. Criteria for inclusion in this figure were (1) the marker corresponds to a signaling receptor such as a growth factor, hormone, or cytokine; and (2) a p value for differential expression below 0.05.

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FIG. 26 is a table listing CNS markers differentially expressed at least in one time point studied in all tumors analyzed.

#### **DETAILED DESCRIPTION**

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The methods described herein rely, in part, on the detection of gene expression in the CNS to identify (e.g., diagnose or monitor) peripheral (non-CNS) tissues or organs for early stages of disease (e.g., in some cases, within hours, days, weeks or months of the appearance of disease). Early identification and/or diagnosis of disease provides an opportunity for early therapeutic intervention to target the disorder before it becomes overly advanced or aggressive.

#### General Methodology

The CNS is involved in the body's response to any internal or external stimulus that by its intensity or functional relevance could alter internal homeostasis. As part of this function, the CNS and the immune system interact to obtain a suitable immune response when necessary.

An immune response impacts the brain via neural and humoral mechanims. Neural mechanisms primarily involve the activation of the vagal nerve. Humoral mechanisms can include cytokine-mediated action directly on brain structures, e.g., cytokine-mediated increases on neural firing rates (Rothwell and Hopkins, 1995, Trends Neurosci 18(3):130-6; Wang et al., 2003, Nature, 421(6921):384-8). In one example, peripheral cytokines have been shown to bind and activate the vagal nerve, which in turn activates neurons of the nucleus of the tractus solitarius and the hypothalamus in the brain (Watkins and Maier, 1999, Proc. Natl. Acad. Sci. USA, 96(14):7710-3).

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Humoral signals from the periphery act as potent messengers to the brain. Cytokines in the brain can exert their action at a much lower dose than in the periphery. For example, intracerebral administration of interleukin-1 (IL-1) at a dose of 100 pg to 10 ng elicits maximal changes in fever, gastric function, increased metabolism and behavioral changes, while several micrograms of this cytokine are necessary to elicit similar responses when administered to the periphery (Rothwell and Hopkins, *supra*).

After sensing an internal immune signal, the brain reacts in different ways. A paradigm of CNS response to immune signals is the activation of neuroendocrine axes such as the hypothalamus-pituitary-adrenal axis. The activation of this axis results in the liberation of glucocorticoids, which in turn can modulate the ongoing immune response in under 10 minutes. Vagatomy has been shown to blunt the activation of the HPA axis after intraperitoneal administration of cytokines (Watkins and Maier, *supra*). This feedback mechanism is of high physiological relevance; i.e., inhibition of glucocorticoid production after cytokine release in the periphery usually results in the death of the organism (Besedovsky and del Rey, 1996, Endocr. Rev., 17(1):64-102).

The brain can also sense signals that will affect the immune and other systems from the external milleu. For example, the triggering of a stress reaction can result in the release of glucocorticoids and the attenuation of an ongoing immune response. The effects of stress on the immune system are well documented in animal models and humans (Deinzer et al., 2000, Int. J. Psychophysiol., 37(3):219-32; Marshall et al., 1998, Brain Behav. Immun., 12(4):297-307; Benschop et al., 1996, FASEB J., 10(4):517-24; Sheridan et al., 1998, Ann. N.Y. Acad. Sci., 840:803-8). In addition, there is anecdotal and preliminary evidence that mind/body interventions such as meditation or yoga could

have an influence on the immune system (Cassileth, 1999, CA Cancer J. Clin., 49(6):362-75).

The new methods harness this natural reaction of the CNS as a way to detect peripheral disease at an early stage.

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#### Cancer Development

It is generally accepted that a clinically detectable tumor mass is composed of cells that, although abnormal, evade immune surveillance and resist immune system attack. During the time of neoplastic progression, cells are characterized by high mutation rates, reflected, inter alia, in phenotypic changes such as downregulation of histocompatibility antigens. A tumor may thus become resistant to a particular therapeutic by clonal selection and proliferation from the tumor mass of a cell clone having a mutation that allows the cell to resist the given therapeutic. The "natural selection" of tumor cell clones occurs at a given rate leading to the appearance of malignant cells having genetic and epigenetic traits that facilitate growth and escape from the immune system. It is estimated that the average malignancy contains more than 10,000 mutations (Stoler et al., 1999, Proc. Natl. Acad. Sci., USA., 96(26):15121-6). Therefore, it can be concluded that the antigen profile of established cancers by no means reflects the cell genotype and phenotype of very early stage neoplasia. Moreover, it is reasonable to assume that tumor antigens present in the established cancer and the response they can induce in the organism will be different than the antigens and responses induced by early stage neoplastic cells. The new methods can detect such early stage neoplastic cells in spite of these obstacles.

Some neoplasms, e.g., some cancers (e.g., certain types of carcinoma) can grow for long periods (e.g., for 1, 2, 5, 10, 15, 20 or 25 years) before they are clinically detectable using prior known technology and/or before they become malignant. This period provides an extraordinary window of opportunity for detection of cancerous cells before the malignant tumor is clinically detectable by current strategies. During this period tumor cells undergo several modifications at the molecular level as a result of their genomic instability.

Each genetic change is potentially selective for proliferation and/or is capable of triggering a new "alarm signal" to recruit and activate local innate and adaptive immune responses. In a simple view, 10,000 alarm signals might be produced during the 10 to 15 years of tumor development before the tumor is clinically detectable.

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#### Development of Rheumatoid Arthritis

Rheumatoid arthritis (RA) is an acquired autoimmune disease in which genetic factors appear to play a role. RA occurs in 1-2 percent of the general population and is found world-wide. Females with RA outnumber males by 3:1. Onset of the disease in adults is usually between the ages of 40 to 60 years, although it can occur at any age.

RA involves Th1 lymphocytes and macrophage infiltration into joints as well as the presence of rheumatoid factors in patients' serum (Chernajovsky et al., 2000, Genes Immun., 1:295-307). Degradation of cartilage is accompanied by the outgrowth of synovial membrane (pannus). This process is generally regulated by IL-1 and TNF-α, while TGF-β and IL-10 counteract this effect (Chernajovsky et al., *ibid*). Susceptibility to arthritis has been correlated with MHC class II locus, in particular HLA-DR4 in 70 percent of patients with RA (Chernajovsky et al., *ibid*). Rheumatoid Factor(s) (RF) are antibodies to IgG, and are present in 60-80 percent of adults with the disease. High titers of RF are usually associated with more severe and active joint disease, greater systemic involvement, and a poorer prognosis for remission.

An unknown antigen is thought to initiate the autoimmune response resulting in RA. It has been suggested that there is a synovial antigen resembling a bacterial lipopolysaccharide (LPS) of arthritogenic bacteria that initiates the autoimmune response (Kennedy, 2000, Med. Hypotheses, 54(5):723-5). TNF- $\alpha$  appears to be the driving force behind the chronic inflammation characteristic of RA. TNF- $\alpha$  plays also an important role in B cell maturation which appears to participate in disease progression (Chernajovsky et al., *ibid*). Some data also strongly indicate a role for Suppressor of Cytokine signaling (SOCS) in disease outcome (Egan et al., 2003, J. Clin. Invest. 111(6):915-24).

The initiation of the autoimmune response and/or the initiation of the inflammatory mechanisms in the early development of RA are likely to trigger signals detected by changes in gene expression in the CNS.

# 5 Development of Asthma

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Asthma is an inflammatory airway disease characterized by the presence of cells such as eosinophils, mast cells, basophils, and CD25+ T lymphocytes in the airway walls. Chemokines attract cells to the site of inflammation and cytokines (Interleukin (IL)-4, IL-5, IL-10 and IL-13) activate them, resulting in inflammation and damage to the mucosa. When asthma becomes chronic, secondary changes occur, such as thickening of basement membrane and fibrosis. IL-4 and other cytokines such as  $TGF-\beta$  may be involved in tissue remodeling and the fibrotic response.

In allergic asthma (also known as extrinsic asthma), the initiation event of airway inflammation is an immunological reaction to allergen. Continued exposure to allergen results in chronic inflammation. Allergic asthma affects about 3 million children (8 to 12 percent of all children) and 7 million adults in the United States at a cost estimated at \$6.2 billion a year. It has been suggested that longitudinal studies based on yet unidentified inflammatory markers will guide asthma management in the future (Wilson, 2002, Curr. Opin. Pulm. Med., 8(1):25-32).

In the development of asthma, the initiation of the allergic or inflammatory response, e.g., release of cytokines and/or chemokines, can likely trigger signals detected by changes in gene expression in the CNS.

## **Development of Obesity**

Body size and body weight are highly heritable traits. Association studies performed with populations of monozygotic and dizygotic twins, non-twin siblings and adoptive family members indicated that the variance for body mass index (body weight divided by height to the square) is much lower in identical twins that in any other group, indicating that genetic factors rather than environmental effects are the key determinant of human adiposity (Maes et al., 1997, Behav. Genet., 27:325-351; Allison et al., 1996, Int. J. Obes. Relat. Metab. Disord. 20:501-506). Diet-induced obesity is also highly

heritable. A pioneer study performed in 12 pairs of young adult identical twins overfed by 1,000 kcal per day during a 100-day period demonstrated that overfeeding induced a variable increase in body weight in all volunteers. However, twin pairs had six times less variance in mass increase than non-twin pairs, indicating that adaptation to long-term overfeeding has important genetic factors (Bouchard et al., 1990, N. Engl. J. Med. 322:1477-1482). The strong genetic predisposition to gain weight after ingesting a fatrich diet is even more clearly observed in the laboratory when testing mice or rats of different genetic backgrounds (Schaffhauser et al., 2002, Obes. Res. 10:1188-1196). Most strains of mice maintain their body weight throughout relatively long periods of time while being fed *ad libitum* with low fat diets. However, when fed *ad libitum* with a high fat diet, some strains develop a considerable increase in body mass and some other strains are resistant to this increase regardless of increase in food consumption (West et al., 1995, Am. J. Physiol., 268:R658-R665; Prpic et al., 2003, Endocrinology, 144:1155-1163).

The regulation of body weight involves a large number of interconnected peripheral and brain circuits that participate in the control of energy balance throughout the entire organism (Spiegelman and Flier, 2001, Cell, 104:531-43). Information about the amount of energy stored in the whole body is transported into the brain by peripheral hormones such as leptin and insulin. The relative variation of the plasma concentration of these hormones is interpreted by central mechanisms to induce signals of appetite or satiety (Friedman and Halaas, 1998, Nature, 395:763-70). Other molecules such as ghrelin and cholecystokinin (CCK) enter into the brain after being released from different portions of the gastrointestinal tract and provide essential information to brain centers about the nutritional status of the organism (Murakami et al., 2002, J. Endocrinol., 174:283-288; Sheng and Moran, 2002, Neuropeptides, 36:171-181).

The hypothalamus, a critical brain area for the complicated control of energy homeostasis, integrates a variety of converging signals within a short time frame. In the ventral hypothalamus a group of appetite-inducing neurons expresses the neuropeptide Y (NPY) gene. As leptin levels drop from circulation NPY is released into the paraventricular nucleus of the hypothalamus to induce food intake (Widdowson et al., 1999, Peptides, 20:367-372). A single intracerebroventricular administration of NPY in

mice or rats can dramatically increase food intake for many hours (Zarjevski et al., 1993, Endocrinology, 133:1753-1758). Conversely, another group of neurons located in the arcuate nucleus of the hypothalamus expresses the proopiomelanocortin gene (POMC). These neurons also express the leptin receptor gene. After an excessive intake of fatenriched food, the levels of triglycerides rise, filling peripheral adipocytes with fat stores. This leads to an increase in production of leptin, which is released into the circulation and eventually enters the brain by a selective uptake mechanism (Hileman et al., 2002, Endocrinology, 143:775-783). Leptin stimulates leptin receptors located in POMC neurons, thereby increasing their firing activity (Cowley et al., 2001, Nature, 411:480-484).

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One of the active peptides produced by the POMC precursor is  $\alpha$ -melanocyte stimulating hormone ( $\alpha$ -MSH). Upon stimulation of leptin receptors,  $\alpha$ -MSH is released in the paraventricular nucleus of the hypothalamus to induce satiety. Intracerebroventricular injections of  $\alpha$ -MSH in mice or rats induce long lasting anorexia that can promote the death of the animals if they are not forced to feed (Fan et al., 1997, Nature, 385:165-168).

The hormones, neuropeptides and their receptors described above are only a few examples of the many gene products that participate in the central control of energy balance. Regulation of a molecule involved in energy control (e.g., a disruption associated with propensity or presence of obesity) can likely trigger signals that result in changes in gene expression in the CNS.

While not limited by any theory, the methods described herein are based, in part, on the discovery that the CNS senses the presence of "alarm signals" from peripheral (non-CNS) disorders at an early stage in the development of disease progression. Thus, the methods described herein relate to diagnosing peripheral disorders by detecting gene expression in the CNS, e.g., in a CNS sample from a subject, such as a human. In one aspect, a non-CNS disorder can be identified based on a profile of gene expression in the CNS (e.g., the brain) within hours, weeks or months after disease progression is initiated in the body. In some embodiments, a non-CNS disorder can be identified based on a profile of gene expression in the CNS (e.g., the brain) within one or more years (e.g., 2,

3, 5, 7, 10 or more years) after disease progression is initiated in the body, but before a disorder is clinically detectable and/or in an advanced stage.

# Methods Of Detecting Gene Expression

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Gene expression in the CNS can be detected in vitro, e.g., in an isolated CNS sample, or in vivo, e.g., using in vivo imaging techniques.

# Central Nervous System (CNS) Samples

The CNS refers to the brain (including the cranial nerves) and spinal cord. A CNS sample can be, e.g., a cell or tissue from the brain or spinal cord, or a sample of the cerebrospinal fluid (CSF) that fills the ventricles of the brain and the central canal of the spinal cord.

Where the detection of gene expression is to be done in a CNS sample isolated from the subject, a CNS sample can be obtained by any number of methods available to the skilled artisan. For example, a CNS cell or tissue sample can be obtained from the brain, e.g., by needle biopsy or by open surgical incision. Imaging of the brain can be performed to determine the precise positioning of the needle or scalpel to enter the brain.

In one example, known as stereotactic biopsy, a tiny hole is drilled into the skull with the patient under light sedation or general anesthesia, and a needle is inserted into the brain tissue guided by computer-assisted imaging techniques such as computerized tomography (CT) or magnetic resonance imaging (MRI) scans. The needle is used to remove a sample of cells, whose gene expression can then be detected by a routine assay, e.g., a gene expression assay described herein. In another example, a sample of CSF can be obtained by routine methods, such as by lumbar puncture. This procedure can be done on an outpatient basis, e.g., under local anesthetic.

The number of cells or amount of CSF needed to perform a particular gene expression assay on a CNS sample will vary; however, some techniques, such as PCR based techniques, will require a very small number of cells, e.g., as few as 10 to 100 cells (Klein et al., Nat. Biotechnol., 20(4):387-92, 2002). The CNS sample can be used immediately in a diagnostic test described herein, or it can be stored, e.g., cooled or frozen, and/or transported to a facility where the diagnostic test is performed.

# Nucleic Acid-Based Methods

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In one embodiment, the methods described herein will utilize techniques for detection of gene expression where a polynucleotide (such as an RNA, mRNA, DNA, cDNA, or other nucleic acid corresponding to the gene) is detected. It should be understood by the skilled artisan that many methods for nucleic-acid based detection of gene expression exist and that any suitable method for detection can be used. Typical assay formats utilize nucleic acid hybridization and include, e.g., 1) nuclear run-on assay, 2) slot blot assay, 3) northern blot assay, 4) magnetic particle separation, 5) nucleic acid or DNA arrays or chips (also discussed in more detail below), 6) reverse northern blot assay, 7) dot blot assay, 8) in situ hybridization, 9) RNase protection assay, 10) ligase chain reaction, 11) polymerase chain reaction (PCR), 12) reverse transcriptase (RT)-PCR, and 13) differential display RT-PCR (DDRT-PCR) or any combination of any two or more of these methods. Such assays can employ the use detectable labels such as radioactive labels, enzyme labels, chemiluminescent labels, fluorescent labels, or other suitable labels, to detect, identify, or monitor the presence or level of a particular nucleic acid being detected. Such techniques and labels are known in the art and widely available to the skilled artisan.

In an exemplary embodiment, an RNase protection assay can be utilized in the methods described herein by hybridizing multiple DNA probes corresponding to one or more members of a panel of sequences to mRNA isolated from a CNS sample from a subject to be tested. The expression profile for one or more genes from the CNS sample can be compared to a reference profile, e.g., a basal pattern of expression, or other negative or positive control (e.g., a profile from a patient known to have no peripheral disease, or a standard or average profile derived from subjects known to not have the particular disorder being tested). In one example, the gene expression profile from the test CNS sample is compared to a reference gene expression profile that is associated with the presence of a non-CNS neoplasia. If the test gene expression profile matches the reference gene expression profile, it indicates that the subject has, or is at risk for developing, the non-CNS neoplastic disorder.

The methods described herein are also well suited for polymerase chain reaction (PCR)-based methods. PCR-based methods include RT-PCR (U.S. Patent No. 4,683,202), ligase chain reaction (Barany, Proc. Natl. Acad. Sci. USA, 88:189-193, 1991), self-sustained sequence replication (Guatelli et al., Proc. Natl. Acad. Sci. USA, 87:1874-1878, 1990), transcriptional amplification system (Kwoh et al., Proc. Natl. Acad. Sci. USA, 86:1173-1177, 1989), Q-Beta Replicase (Lizardi et al., BioTechnology, 6:1197, 1988), rolling circle replication (Lizardi et al., U.S. Patent No. 5,854,033), or any other nucleic acid amplification method, followed by the detection of the amplified molecules using techniques known in the art. PCR amplification of mRNAs expressed in a CNS sample can be performed directly from mRNA isolated from the sample, or from cDNA reverse-transcribed from such isolated mRNA. The amplified nucleic acid can then be hybridized to a particular probe of interest, e.g., a probe for a CNS gene as described herein, in order to determine its expression. The probe can be disposed on an address of an array, e.g., an array described herein below. Such methods are routine and are particularly amendable to routine adaptation to automated systems employing computer controlled reagent aliquoting and signal detection. See, e.g., Klein et al., Nat. Biotechnol., 2002, 20(4):387-92.

In another embodiment, in situ methods are used to detect the presence or level of mRNA corresponding to a particular gene. In such methods, a CNS cell or tissue sample can be prepared/processed and immobilized on a support, typically a glass slide, and then contacted with a probe (e.g., a probe for a CNS gene described herein).

In still another embodiment, serial analysis of gene expression, as described in U.S. Patent No. 5,695,937, is used to detect transcript levels of a CNS gene described herein.

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### Polypeptide-Based Methods

In one embodiment, the methods described herein utilize techniques for detection of gene expression where a gene product (polypeptide) encoded by a gene is detected or where an activity of the polypeptide, e.g., an enzymatic activity, is detected. Such methods are particularly advantageous for detecting the expression of genes that encode polypeptides that are secreted from CNS cells, e.g., into the CSF.

A variety of methods can be used to determine the level of protein encoded by a CNS gene. In general, these methods include contacting a CNS sample (such as a brain cell sample or a CSF sample) with an agent, such as an antibody, that selectively binds to the protein of interest. In one embodiment, the antibody bears a detectable label.

Antibodies can be polyclonal, or more preferably, monoclonal. An intact antibody, or a fragment thereof (e.g., Fab or F(ab')2) can be used. The term "labeled," with regard to the probe or antibody, is intended to encompass direct labeling of the probe or antibody by coupling (i.e., physically linking) a detectable substance to the probe or antibody, as well as indirect labeling of the probe or antibody by reactivity with a detectable substance. Such detection methods can be used to detect a CNS gene product in a CNS sample in vitro as well as in vivo.

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In vitro techniques include immunoassays such as enzyme linked immunosorbent assays (ELISAs), immunoprecipitations, immunofluorescence, enzyme immunoassay (EIA), radioimmunoassay (RIA), Western blot analysis, and Luminex<sup>TM</sup> x MAP<sup>TM</sup> detection assay. Some immunoassays are "sandwich" type assays, in which a target analyte(s) is "sandwiched" between a labeled antibody and an antibody immobilized onto a solid support. The assay is read by observing the presence and amount of antigenlabeled antibody complex bound to the immobilized antibody. Another immunoassay useful in the methods described herein is a "competition" type immunoassay, wherein an antibody bound to a solid surface is contacted with a sample (e.g., a CSF sample) containing both an unknown quantity of antigen analyte and with labeled antigen of the same type. The amount of labeled antigen bound on the solid surface is then determined to provide an indirect measure of the amount of antigen analyte in the sample. Such immunoassays are readily performed in a "dipstick" format (e.g., a flow-through or migratory dipstick design) for convenient use. A dipstick-based assay optionally includes an internal negative or positive control. Numerous types of dipstick immunoassays assays are known in the art and are described, e.g., in U.S. Patents 5,656,448; 4,366,241; and 4,770,853. In other embodiments, antibody based assays are performed in an array format. For example, a CNS sample is labeled, e.g., biotinylated, and then contacted to an antibody, e.g., an antibody positioned on an antibody array. The sample can be detected, e.g., with avidin coupled to a fluorescent label.

In vivo techniques include, e.g., introducing into a subject (e.g., into the CSF) a labeled antibody that binds to the gene product to be detected. The antibody can be labeled, e.g., with a radioactive marker, whose presence and location in a subject can be detected by standard imaging techniques.

Polyclonal and monoclonal antibodies to be used to detect a particular CNS gene product will, in certain cases, be available. For example, commercially available antibodies exist for many of the CNS marker genes described herein. Alternatively, a skilled artisan can make a suitable antibody for use in a diagnostic assay using routine techniques. Methods of making and using polyclonal and monoclonal antibodies to detect a particular target are described, e.g., in Harlow et al., <u>Using Antibodies: A Laboratory Manual: Portable Protocol I.</u> Cold Spring Harbor Laboratory (December 1, 1998). Methods for making modified antibodies and antibody fragments (e.g., chimeric antibodies, reshaped antibodies, humanized antibodies, or fragments thereof, e.g., Fab', Fab, F(ab')2 fragments); or biosynthetic antibodies (e.g., single chain antibodies, single domain antibodies (DABs), Fv, single chain Fv (scFv), and the like), are known in the art and can be found, e.g., in Zola, <u>Monoclonal Antibodies: Preparation and Use of Monoclonal Antibodies and Engineered Antibody Derivatives</u>, Springer Verlag (December 15, 2000; 1st edition).

## 20 <u>Imaging of CNS Gene Expression</u>

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In one embodiment, the methods described herein utilize techniques for imaging of gene expression, e.g., non-invasive imaging of gene expression, in the CNS. For example, a labeled probe that is capable of detecting the expression of a target gene can be delivered into the brain through the blood-brain barrier (BBB) by targeting the labeled probe to the brain via endogenous BBB transport systems, such as carrier-mediated transport systems that exist for the transport of nutrients across the BBB. Similarly, receptor-mediated transcytosis systems operate to transport circulating peptides across the BBB, such as insulin, transferrin, or insulin-like growth factors. These endogenous peptides can act as "transporting peptides," or "molecular Trojan horses," to ferry a labeled diagnostic probe as described herein, across the BBB. The label can then be detected by known brain imaging techniques. Such an approach is described, e.g., in

U.S. Patent No. 6,372,250. In other embodiments, Shi et al., Proc. Natl. Acad. Sci. USA, 2000, 97(26):14709-14 and Lee et al. J. Nucl. Med. 2002, 43(7):948-56 describe imaging of gene expression in the brain *in vivo* using an antisense radiopharmaceutical combined with drug-targeting technology to traverse the BBB.

Other methods of delivering into the brain a labeled probe that is capable of detecting the expression of a target gene are described, e.g., in U.S. Pat. No. 5,720,720. This patent describes methods of delivering agents (such as labeled antibodies for imaging gene products) into the brain by high-flow microinfusion.

# Detection of Changes in CNS Gene Expression in Bodily Fluids

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In some cases, gene activation in the CNS can result in a measurable alteration in a gene product at a distant site, e.g., in a fluid such as blood, urine or semen. It is known, e.g., that the cerebral cortex, hippocampus, entorrhinal cortex, parts of the thalamus, basal ganglia, cerebellum and the reticular formation influence the output of the autonomic nervous system (Kandel et al, Principles of Neural Science, Third Edition, Appleton & Lange). These influences could result in measurable alterations of gene expression at the mRNA or protein level in autonomic ganglia or in inervated organs. An example of this type of interaction is the immunomodulatory action of the activation of the vagus nerve after cytokine release in the periphery (Tracey, Nature, 420:853-9, 2002).

In addition, gene activation in the CNS can be detected by measuring changes in blood proteins in some cases. For example, neurons in the CNS can trigger the release of hormones in blood via the activation of several neuroendocrine axes such as the hypothalamus-pituitary-adrenal, -gonadal or thyroid axes (Besedovsky and del Rey, Endocrine Reviews, 17:1-39, 1996). Moreover, brain extracellular fluid could drain into blood and deep cervical lymph (Cserr et al, Brain Pathol., 2(4):269-76, 1992). Cerebral extracellular fluids drain from brain to blood across the arachnoid villi and to lymph along certain cranial nerves (primarily olfactory) and spinal nerve root ganglia. A minimum of 14 to 47% of protein injected into different regions of brain or cerebrospinal fluid passes through lymph. Thus, CSF markers could drain into, and be detected in, lymph, blood or serum. Such markers found in blood may also be enriched, and thereby detectable, in urine, due to selective filtration of blood components by the kidneys.

The CNS is connected to the testis via the autonomic nervous system as well as the endocrine system. If a change in gene activity in the brain results in modifications in the activity of the hypothalamus-pituitary-gonadal axis or in the innervation of the testes, these changes could be then detected in fluids related to the testes, such as semen. For example, patients with spinal cord injury have been shown to have alterations in the composition of their semen (See Naderi and Safarinejad, Clin. Endocrinol., 58(2):177-84, 2003).

Routine methods can be used to identify gene products in peripheral tissues, such as peripheral bodily fluids, which are the result of changes in gene expression in the CNS. For example, a candidate marker gene can be disrupted in the brain of an experimental animal. A change in the expression of a candidate gene in a peripheral tissue in the experimental animal, compared to a wild type animal, indicates that the expression of the candidate molecule in the peripheral tissue is tied to changes in gene expression in the CNS.

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#### <u>Arrays</u>

The methods described herein are readily adapted for nucleic acid or protein arrays, e.g., nucleic acid and/or protein "chips," following the methods and teachings known in the art. In a typical embodiment, an array chip includes multiple probes (e.g., DNA probes and/or antibody probes) for detection of expression of multiple CNS genes. In one embodiment, the probes on a specific chip are chosen to detect the members of one or more specific panels or "clusters" of genes, each cluster being associated with a specific gene expression profile if a non-CNS neoplasia is present in the subject from whom the CNS sample was taken. A chip can contain tens, hundreds, or thousands of individual probes immobilized (tethered) at discrete, predetermined locations (addresses or "spots") on a solid, planar support, e.g., glass, metal, or nylon. An array can be a macroarray or microarray, the difference being in the size of the spots. Macroarrays contain spots of about 300 microns in diameter or larger and can be imaged using gel or blot scanners. Microarrays contain spots less than 300 microns, typically less than 200 microns, in diameter.

For analysis and comparison of profiles of gene expression in the methods described herein, a nucleic acid array can be constructed using nucleic acid probes for at least four, e.g., at least 10, 20, 40, 60, 80 or 100 CNS genes. Such an array can include control probes (i.e., probes for genes whose expression is expected to remain unaffected in a negative sample, e.g., a sample from a subject not having a non-CNS disorder). Typically, such controls or "normal" non-disease samples are obtained from healthy volunteers. Longitudinal studies of healthy volunteers can be performed to confirm that the control samples are from individuals that remained disease free. Such studies can provide the raw data for a database of control gene expression profiles. Such a database can provide a source of normal or control "reference" profiles that can be used in the present methods. Control samples can also be obtained post-mortem from individuals who died for a reason unrelated to the disorder being diagnosed (e.g., individuals who died from an accidental trauma). In such cases, post-mortem samples should be taken as soon as possible after death, e.g., no later than 3 hours after death.

A population of labeled cDNA representing total mRNA from a sample of a tissue of interest, e.g., brain, spinal cord or CSF, is contacted with the DNA array under suitable hybridization conditions. Hybridization of cDNAs with sequences in the array is detected, e.g., by fluorescence at particular addresses on the solid support. Thus, a pattern of fluorescence representing a gene expression pattern in the CNS sample of a particular subject or group of subjects is obtained. These patterns of gene expression can be digitized and stored electronically for computerized analysis and comparison. For example, an array can be used to compare expression of CNS genes in individuals being tested with one or more reference gene expression profiles stored electronically, e.g., in a digital database, where the reference gene expression profile is associated with either the presence or absence of a peripheral neoplasia.

In some embodiments, cDNAs are used as probes to form the array. Suitable cDNAs can be obtained by conventional polymerase chain reaction (PCR) techniques, as described above. The length of the cDNAs can be from 20 to 2,000 nucleotides, e.g., from 100 to 1,000 nucleotides. Other methods known in the art for producing cDNAs can be used. For example, reverse transcription of a cloned sequence can be used (for example, as described in Sambrook et al., eds., Molecular Cloning: A Laboratory

Manual. 2nd ed., Cold Spring Harbor Laboratory, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY, 1989). The cDNA probes are deposited or placed ("printed" or "spotted") onto a suitable solid support (substrate), e.g., a coated glass microscope slide, at specific, predetermined locations (addresses) in a two-dimensional grid. A small volume, e.g., 5 nanoliters, of a concentrated DNA solution is used in each spot. Spotting can be carried out using a commercial microspotting device (sometimes called an arraying machine or gridding robot) according to the vendor's instructions. Commercial vendors of solid supports and equipment for producing DNA arrays include BioRobotics Ltd., Cambridge, UK; Corning Science Products Division, Acton, MA; GENPAK Inc., Stony Brook, NY; SciMatrix, Inc., Durham, NC; and TeleChem International, Sunnyvale, CA.

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The cDNAs can be attached to the solid support by any suitable method. In general, the linkage is covalent. Suitable methods of covalently linking DNA molecules to the solid support include amino cross-linking and UV crosslinking. For guidance concerning construction of cDNA arrays according to the invention, see, e.g., DeRisi et al., Nature Genetics, 1996, 14:457-460; Khan et al., Electrophoresis, 1999, 20:223-229; Lockhart et al., Nature Biotechnol., 1996, 14:1675-1680.

In some embodiments of the invention, the immobilized DNA probes in the array are synthetic oligonucleotides. Preformed oligonucleotides can be spotted to form a DNA array, using techniques described herein with regard to cDNAs. In general, however, the oligonucleotides are synthesized directly on the solid support. Methods for synthesizing oligonucleotide arrays are known in the art. See, e.g., Fodor et al., U.S. Patent No. 5,744,305. The sequences of the oligonucleotides represent portions of the sequences of a particular gene to be detected above. Generally, the lengths of oligonucleotides are 10 to 50 nucleotides, e.g., 15, 20, 25, 30, 35, 40, or 45 nucleotides.

Also useful in the methods are aptamer arrays. Aptamers are nucleic acid molecules that bind to specific target molecules based on their three-dimensional conformation rather than hybridization. The aptamers are selected, for example, by synthesizing an initial heterogeneous population of oligonucleotides, and then selecting oligonucleotides within the population that bind tightly to a particular target molecule. Once an aptamer that binds to a particular target molecule has been identified, it can be

replicated using a variety of techniques known in biological and other arts, e.g., by cloning and polymerase chain reaction (PCR) amplification followed by transcription. The target molecules can be nucleic acids, proteins, peptides, small organic and inorganic compounds, and even entire micro-organisms.

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The synthesis of a heterogeneous population of oligonucleotides and the selection of aptamers within that population can be accomplished using a procedure known as the Systematic Evolution of Ligands by Exponential Enrichment or SELEX. The SELEX method is described in, e.g., Gold et al., U.S. Patent Nos. 5,270,163 and 5,567,588; Fitzwater et al., ("A SELEX Primer," Methods in Enzymology, 267:275-301, 1996); and in Ellington and Szostak ("In Vitro Selection of RNA Molecules that Bind Specific Ligands," Nature, 346:818-22). Briefly, a heterogeneous DNA oligomer population is synthesized to provide candidate oligomers for the in vitro selection of aptamers. This initial DNA oligomer population is a set of random sequences 15 to 100 nucleotides in length flanked by fixed 5' and 3' sequences 10 to 50 nucleotides in length. The fixed regions provide sites for PCR primer hybridization and, in one implementation, for initiation of transcription by an RNA polymerase to produce a population of RNA oligomers. The fixed regions also contain restriction sites for cloning selected aptamers. Many examples of fixed regions can be used in aptamer evolution. See, e.g., Conrad et al. ("In Vitro Selection of Nucleic Acid Aptamers That Bind Proteins," Methods in Enzymology, 267:336-83, 1996); Ciesiolka et al., ("Affinity Selection-Amplification from Randomized Ribooligonucleotide Pools," Methods in Enzymology, 267:315-35, 1996); Fitzwater, supra.

Aptamers are generally selected in a 5 to 100 cycle procedure. In each cycle, oligomers are bound to the target molecule, purified by isolating the target to which they are bound, released from the target, and then replicated by 20 to 30 generations of PCR amplification.

Aptamer selection is similar to evolutionary selection of a function in biology. Subjecting the heterogeneous oligonucleotide population to the aptamer selection procedure described above is analogous to subjecting a continuously reproducing biological population to 10 to 20 severe selection events for the function, with each selection separated by 20 to 30 generations of replication.

Heterogeneity is introduced, e.g., only at the beginning of the aptamer selection procedure, and does not occur throughout the replication process. Alternatively, heterogeneity can be introduced at later stages of the aptamer selection procedure.

Various oligomers can be used for aptamer selection, including, e.g., 2'-fluororibonucleotide oligomers, NH2-substituted and OCH3-substituted ribose aptamers, and
deoxyribose aptamers. RNA and DNA populations are equally capable of providing
aptamers configured to bind to any type of target molecule. Within either population, the
selected aptamers occur at a frequency of 109 to 1013, see Gold et al., ("Diversity of
Oligonucleotide Functions," Annual Review of Biochemistry, 64:763-97, 1995), and most
frequently have nanomolar binding affinities to the target, affinities as strong as those of
antibodies to cognate antigens. See Griffiths et al., (EMBO J., 13:3245-60, 1994).

Using 2'-fluoro-ribonucleotide oligomers is likely to increase binding affinities ten to one hundred fold over those obtained with unsubstituted ribo- or deoxyribo-oligonucleotides. See Pagratis et al. ("Potent 2'-amino and 2' fluoro 2'deoxyribonucleotide RNA inhibitors of keratinocyte growth factor" Nature Biotechnology, 15:68-73). Such modified bases provide additional binding interactions and increase the stability of aptamer secondary structures. These modifications also make the aptamers resistant to nucleases, a significant advantage for real world applications of the system. See Lin et al. ("Modified RNA sequence pools for in vitro selection" Nucleic Acids Research, 22:5229-34, 1994); Pagratis, *supra*.

In the present invention, aptamers can be used to detect, e.g., mRNAs, cDNAs, or proteins corresponding to CNS marker genes.

In some embodiments of the invention, probes (e.g., nucleic acid probes, antibodies, or aptamers) for the human homologs of CNS genes are used in the detection method. In other embodiments, the probe used for detection consists of highly conserved regions of a gene, e.g., a sequence that is highly conserved between homologous mouse and human sequence.

# Sample Preparation and Analysis

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In methods of the invention, the transcription level of one or more CNS genes is assumed to be reflected in the amount of its corresponding mRNA present in cells of an

assayed CNS sample. In general, mRNA from the CNS cells or tissue is copied into cDNA under conditions such that the relative amounts of cDNA produced representing specific genes reflect the relative amounts of the mRNA in the sample. Comparative hybridization methods involve comparing the amounts of various, specific mRNAs in two tissue samples, as indicated by the amounts of corresponding cDNAs hybridized to sequences from the genes of interest.

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The mRNA used to produce cDNA is generally isolated from other cellular contents and components. One useful approach for mRNA isolation is a two-step approach. In the first step, total RNA is isolated. The second step is based on hybridization of the poly(A) tails of mRNAs to oligo(dT) molecules bound to a solid support, e.g., a chromatographic column or magnetic beads. Total RNA isolation and mRNA isolation are known in the art and can be accomplished, for example, using commercial kits according to the vendor's instructions. Similarly, synthesis of cDNA from isolated mRNA is known in the art and can be accomplished using commercial kits according to the vendor's instructions. Fluorescent labeling of cDNA can be achieved by including a fluorescently labeled deoxynucleotide, e.g., Cy5-dUTP or Cy3-dUTP, in the cDNA synthesis reaction. For guidance concerning isolation of mRNA and synthesis of fluorescently labeled cDNA for analysis on a DNA array, see, e.g., Ross et al., Nature Genetics 2000, 24:227-235.

In the invention, conventional techniques for hybridization and washing of DNA arrays, detection of hybridization, and data analysis can be employed routinely without undue experimentation. Commercial vendors of hardware and software for scanning DNA arrays and analyzing data include Cartesian Technologies, Inc. (Irvine, CA); GSI Lumonics (Watertown, MA); Genetic Microsystems Inc. (Woburn, MA); and Scanalytics, Inc. (Fairfax, VA).

In other embodiments, the expression level of one or more CNS genes is reflected in the presence and/or level of protein present in cells of a CNS sample to be assayed. The presence or level of protein in a CNS sample can be detected by routine methods. For example, a CNS sample (e.g., a CSF sample) can be analyzed by gel electrophoresis techniques such as 2-dimensional (2D) PAGE. Once protein spots are separated on a 2D-PAGE gel, differentially expressed spots can be identified, e.g., by matrix assisted laser

desorption ionization time of flight (MALDI-TOF) and electrospray ionization (ESI). This method can also be used for peptide analysis to provide the fingerprint of a particular protein in a sample.

A second proteomic approach can involve obtaining a proteomic spectrum by directly analyzing a CNS sample, such as a CSF sample, by mass spectroscopy. For example, surface enhanced laser desorption ionization time of flight (SELDI-TOF) analysis can be performed to generate a proteomic pattern from a CNS sample. SELDI-TOF analysis has been shown to be able to identify a cluster pattern that differentiates between normal and disease patients. See, Paweletz et al., Dis. Markers, 17(4):301-7, 2001.

## Generating Gene Expression Profiles

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A gene expression profile used in the methods described herein is a pattern of expression of two or more CNS genes. In some cases, an expression profile can be a pattern of expression of 5, 10, 25, 50, 100, 200, 500, or more genes. A "reference gene expression profile" as used herein is a characteristic pattern of expression of two or more CNS genes, where the pattern of expression is associated with risk or presence of a particular disorder. The association between the characteristic profile and the particular disorder is determined through the generation and analysis of CNS gene expression data to mine and identify correlations between particular patterns of CNS gene expression (e.g., relative increases and/or decreases of gene expression of particular genes compared to a negative control) and particular clinical states. For example, a reference gene expression profile can be a set of genes (also referred to herein as a "panel" or "cluster" of genes), where each gene of the set is either down-regulated or upregulated when associated with a specific peripheral disorder or any peripheral disorder. A reference profile can also include a value, e.g., a relative value, of gene expression for two or more genes in a panel, where at least one gene of the panel is down-regulated and at least one gene is up-regulated. An example of such a gene expression profile is a profile that includes a value for the relative differential expression of at least 2, e.g., between 2 and 50, of the genes shown in any of the tables of FIG. 24A-C or between two and seven of the genes listed in FIG. 26. Such a reference profile is associated with the presence of

early stage carcinoma. Other examples are provided by each of the clusters disclosed in FIGS. 2-26. For example, clusters 9 and 10 of FIG. 13(1-5) each provide a profile or panel of genes that are strongly down-regulated in the cortex in response to the presence of lung cancer.

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Exemplary gene expression profiles associated with non-CNS carcinoma (or particular types of non-CNS carcinoma, such as breast, lung or colon carcinoma) are shown in FIGS. 2-26. A reference gene expression profile can include at least a portion of the genes or gene products shown in these figures. For example, a reference gene expression profile associated with lung carcinoma can include a value for the differential expression of 1, 2, 5, 10, 20, 30, 40, 50, or more, genes or gene products listed as CNS markers for lung carcinoma in FIG. 24C. In another example, a reference gene expression profile associated generally with carcinoma can include a value for the differential expression of between one and seven genes or gene products listed as CNS markers for carcinoma in FIG. 26. The reference profiles that can be used with the methods of the invention are not limited by the CNS markers described herein.

Reference profiles can be generated by detecting changes in patterns of gene expression in the CNS in response to the presence of non-CNS disease in an experimental animal, and identifying the human homologs of the genes and gene clusters that are differentially expressed in a certain pattern in the experimental samples, as exemplified in Examples 1-3 described herein. A reference gene expression profile can also be obtained by evaluating human CNS gene expression data. For example, a database can be created and maintained where CNS gene expression data is obtained and stored, e.g., digitally or electronically, for tens, hundreds, or thousands of individuals. The individuals can be followed and evaluated with regard to cancer clinical state longitudinally (e.g., at least 5 years, 10 years, 15 years, 20 years, 30 years, 50 years or a lifetime). The expression profiles of individuals who developed a particular disease, e.g., 5, years, 10 years, 15 years, 20 years, 30 years, or 50 years after the CNS gene expression data was obtained, can be compared with the expression profiles of individuals who remained disease free. Similar comparison can be made between individuals who developed one clinical type of the disorder compared to another, or individuals who developed the disease at an early age versus a late age. These analyses can provide specific reference CNS gene

expression profiles that are associated with different stages of disease, e.g., different stages of neoplasia, or different types of tumors.

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A "test gene expression profile" is obtained from a CNS sample of a subject to be tested for the presence of peripheral disease. First, a CNS sample, e.g., a brain cell sample or CSF sample, is obtained from the subject by routine means such as brain needle biopsy (for a brain cell sample) or a lumbar puncture (for CSF), as described herein. The sample is then prepared for use in a method of detecting gene expression, e.g., any method of detecting gene expression described herein. In one embodiment, total RNA can be prepared from the sample, and reverse transcribed into cDNA for use in a nucleic acid array assay described herein. In another embodiment, total protein is prepared from the sample for use in an antibody assay described herein. The prepared sample can then be contacted with an array (e.g., an antibody or nucleic acid array) that can detect expression levels (or protein levels in the case of an antibody array) of at least one cluster or panel of CNS genes or gene products corresponding to the cluster or panel of CNS genes or gene products of one or more particular reference gene expression profiles to which the test sample will be compared. For example, a prepared CNS sample from the test subject can be contacted with a nucleic acid array containing nucleic acid probes or an antibody array containing antibody probes for two or more, e.g., between 2 and 50, between 2 and 100, or between 10 and 500, of the genes shown in FIGS. 2-26. In one embodiment, the array can contain probes for each of the marker genes in a particular cluster disclosed in any of FIGS. 2-26.

The results of the array assay are obtained by routine techniques, such as fluorescence detection and measurement of bound antibody or hybridized nucleic acid for each position (each probe) on the array. A dataset of the values for the level of each polypeptide or gene detected in the CNS sample by each antibody or probe on the array can then be generated. The dataset can contain information such as patient identifier, and actual and/or relative levels of expression or protein detected. Such a dataset can be used directly as the "test gene expression profile" or the dataset can be converted into a format comparable to the format of the reference profile.

Once the test expression profile is generated, a test profile can be compared to a reference expression profile as described herein.

# **Analyzing Gene Expression Profiles**

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The invention also features methods of evaluating a subject by comparing a test gene expression profile from a test subject with a reference gene expression profile, e.g., a negative control ("normal") gene expression profile associated with the absence of a particular non-CNS disorder or a positive control gene expression profile associated with the presence of the disorder. Longitudinal studies of CNS gene expression in multiple volunteers can be performed to identify and confirm reference gene expression profiles that are associated with individuals who remain disease free or individuals who get the disease. Such studies can provide the raw data for a database of negative and positive control gene expression profiles that can be used in the present methods.

Subject "test" and "reference" profiles can be obtained by methods described herein. In one embodiment, the method includes obtaining a CNS sample from a subject (either directly or indirectly from a caregiver or other party), creating an expression profile from the sample, and comparing the subject's expression profile to one or more reference profiles and/or selecting a reference profile most similar to that of the subject.

As with other detection methods, profile-based assays can be performed prior to the onset of symptoms (in which case they can be diagnostic), prior to treatment (in which case they can be prognostic) or during the course of treatment (in which case they serve as monitors) (see, e.g., Golub et al., 1999, Science 286:531).

A variety of routine statistical measures can be used to compare two gene expression profiles. One possible metric is the length of the distance vector that is the difference between the two profiles. Each of the test and reference profile is represented as a multi-dimensional vector, wherein each dimension is a value in the profile, e.g., a value for the expression of a particular gene in a panel. A test profile and reference profile can be said to match if they are at least 70% identical in reference to the number of genes having similar expression patterns in each profile, or to the level of expression of the genes in each profile. In one embodiment, a test and reference profile are said to match if their respective multi-dimensional vectors, as described above, have a 30% or lower variance with respect to each other. If the test and reference profile match, the test subject can be identified as having the peripheral disorder with which the reference

profile is associated. If the test and normal profile match, the subject is likely to be free of the peripheral disorder.

In one embodiment, pattern recognition software is used to identify matching profiles. For example, unsupervised clustering algorithms, such as hierarchical clustering, K-means clustering, and SOM (self-organizing maps) for pattern discovery can be used. Supervised techniques such as SVM (support vector machines) and SPLASH (structural pattern localization analysis by sequential histograms) algorithms implemented in the Genes@Work software package (IBM Corp.) can also be used.

In another embodiment, gene expression profiles are analyzed by quantitative pattern comparison performed by applying a nearest neighbor classifier (see Jelinek et al., Mol. Cancer Res., 1:346-61, 2003). Based on the nearest neighbor classifier a score is defined which, together with a permutations-derived distribution, can be used to estimate the probability of each test profile of belonging to a class defined by a reference gene expression pattern (see Jelinek, *supra*).

The result of the diagnostic test, which can be transmitted to the subject, a caregiver, or another interested party, can be the subject expression profile per se, a result of a comparison of the subject expression profile with another profile, a most similar reference profile, or a descriptor of any of these. Transmission can occur across a computer network (e.g., in the form of a computer transmission such as a computer data signal embedded in a carrier wave). Accordingly, the invention also features a computer medium having executable code for effecting the following steps: receive a subject expression profile; access a database of reference expression profiles; and either i) select a matching reference profile most similar to the subject expression profile, or ii) determine at least one comparison score for the similarity of the subject expression profile to at least one reference profile. The subject expression profile and the reference expression profile each include a value representing the level of expression of one or more of the identified genes or gene products or the proteins they encode.

### Predictive Medicine

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The methods described herein are generally useful in the field of predictive medicine and, more specifically, are useful in diagnostic and prognostic assays, in

monitoring progression of a disease, e.g., neoplasia, or monitoring of response to treatment, e.g., in clinical trials. For example, one can determine whether a subject has a very early stage neoplasia, in the absence of other, e.g., clinical, indications of neoplasia. In another example, one can determine whether a subject is at risk for developing rheumatoid arthritis or whether the subject has early stage RA, in the absence of clinical indications of RA such as joint inflammation. The methods are particularly useful, e.g., for patients who have had surgery or treatment for the disease (e.g., to remove cancer), in which case the methods could be used to monitor recurrence or metastasis, for persons living in regions of high incidence of cancer due, e.g., to environmental factors, or for individuals who have a family history of a disease (e.g., diabetes, asthma or cancer) or are carriers of a disease susceptibility gene, e.g., a cancer susceptibility gene (e.g., BRCA1 or BRCA2, hMSH2, MLH1, MSH2, or MSH6). Other cancer susceptibility genes are described in The Genetic Basis of Human Cancer, 2nd edition (Vogelstein and Kinzler, Eds.), McGraw-Hill Professional (2002). Such individuals can be evaluated using the methods described herein.

In some cases, for example, where the risk of developing a disease is high (e.g., where an individual has a strong family history of asthma, or where an individual carries a cancer susceptibility gene or lives in a high risk area for cancer), an individual an be evaluated periodically (e.g., every 10 years, every 5 years, or every year) during his lifetime.

The "subject" referred to here, and that is referred to in the context of any of the methods of the invention, is a vertebrate animal, typically a mammal. The subject can be an experimental animal (e.g., an experimental rodent such as a rat or mouse), a domesticated animal (e.g., a dog or cat); an animal kept as livestock (e.g., a pig, cow, sheep, goat, or horse); a non-human primate (e.g., an ape, monkey, or chimpanzee). The animal can be an unborn animal (accordingly, the methods of the invention can be used to carry out genetic screening or to make prenatal diagnoses). Of course, the subject can also be, and typically is, a human.

#### Computer-Readable Medium

In another aspect, the invention features a computer-readable medium having a plurality of digitally encoded data records. Each data record includes a value representing the level of expression of a CNS gene, and a descriptor of the sample. The descriptor can be, e.g., an identifier (e.g., an identifier for the patient from which the sample was obtained, e.g., a name or a reference code that can be matched with patient information only by those having access to a decoding table), a diagnosis made, or a treatment to be performed in the event the level of expression reaches a certain level or falls below a certain level. The data record can also include values representing the level of expression of related genes (e.g., the data record can include values for each of a plurality of genes in a gene "cluster," where a particular reference profile of gene expression for the genes in the cluster is associated with a peripheral disorder). The data record can also include values for control genes (e.g., genes whose expression is not changed in control samples or whose expression is not diagnostically correlated with a peripheral disorder). The data record can be structured as a table (e.g., a table that is part of a database such as a relational database (e.g., a SQL database of the Oracle or Sybase database environments)).

#### Non-CNS Diseases

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The methods described herein are not limiting in that they can be used to diagnose, monitor, or treat any non-CNS disorder, such as a neoplasia (e.g., tumor or cancer); an immune disorder (e.g., an autoimmune disorder such as rheumatoid arthritis, multiple sclerosis, systemic lupus erythematosus, psoriasis, scleroderma); an allergic or inflammatory disorder (e.g., asthma, inflammatory bowel disease, Crohn's disease); a metabolic or endocrine disorder (e.g., diabetes, obesity, Addison's disease); a pathogenic infection (e.g., a viral, parasitic or fungal infection, e.g., HIV infection); or a cardiovascular disorder.

As used herein, "neoplasia" refers to the uncontrolled and progressive proliferation of cells under conditions that would not elicit, or would cause cessation of, proliferation of normal cells. Neoplasia results in the formation of a "neoplasm," which is defined herein to mean any new and abnormal growth, particularly a new growth of

tissue, in which the growth is uncontrolled and progressive. Neoplasm, as used herein, is synonymous with "tumor." Malignant neoplasms or tumors are distinguished from benign in that the former show a greater degree of anaplasia, or loss of differentiation and orientation of cells, and have the properties of invasion and metastasis. Thus, neoplasia includes "cancer," which herein refers to a proliferation of cells having the unique trait of loss of normal controls, resulting in unregulated growth, lack of differentiation, local tissue invasion, and metastasis. The methods described herein can be used to diagnose neoplasia from any non-CNS cell or tissue type, such as neoplasia derived from epithelial or endocrine tissue, mesenchymal tissues, or hematopoietic tissue.

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The term "carcinoma" is art recognized and refers to malignancies of epithelial or endocrine tissues including respiratory system carcinomas, gastrointestinal system carcinomas, genitourinary system carcinomas, testicular carcinomas, breast carcinomas, prostatic carcinomas, endocrine system carcinomas, and melanomas. Exemplary carcinomas include those forming from tissue of the colon, lung, prostate, breast, cervix, head and neck, and ovary. The term also includes carcinosarcomas, which include malignant tumors composed of carcinomatous and sarcomatous tissues. An "adenocarcinoma" refers to a carcinoma derived from glandular tissue or in which the tumor cells form recognizable glandular structures.

The term "sarcoma" is art recognized and refers to malignant tumors of mesenchymal derivation.

As used herein, the term "hematopoietic neoplastic disorders" includes diseases involving hyperplastic/neoplastic cells of hematopoietic origin, e.g., arising from myeloid, lymphoid or erythroid lineages, or precursor cells thereof. The disorders can arise from poorly differentiated acute leukemias, e.g., erythroblastic leukemia and acute megakaryoblastic leukemia. Exemplary myeloid disorders include, but are not limited to, acute promyeloid leukemia (APML), acute myelogenous leukemia (AML) and chronic myelogenous leukemia (CML) (reviewed in Vaickus, L. (1991) Crit Rev. in Oncol./Hemotol. 11:267-97); lymphoid malignancies include, but are not limited to acute lymphoblastic leukemia (ALL) which includes B-lineage ALL and T-lineage ALL, chronic lymphocytic leukemia (CLL), prolymphocytic leukemia (PLL), hairy cell leukemia (HLL) and Waldenstrom's macroglobulinemia (WM). Additional forms of

malignant lymphomas include, but are not limited to non-Hodgkin lymphoma and variants thereof, peripheral T cell lymphomas, adult T cell leukemia/lymphoma (ATL), cutaneous T-cell lymphoma (CTCL), large granular lymphocytic leukemia (LGF), Hodgkin's disease and Reed-Stemberg disease.

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# Identification Of CNS Marker Genes for Non-CNS Disorders

Also featured in the invention are methods of identifying a CNS diagnostic marker for a non-CNS disorder in a subject. Generally, such methods involve detecting changes in gene expression in the CNS in response to the presence of a particular non-CNS disease condition in a subject, e.g., an experimental animal. The methods will generally involve inducing a disease condition or disorder in a test experimental animal; and comparing the expression of at least one gene in a CNS sample from the test experimental animal to expression of the gene in a CNS sample from a control experimental animal. A gene (or a human homolog of a gene) that is differentially expressed in the CNS sample from the test experimental animal compared to the CNS sample from the control experimental animal can be identified as a CNS diagnostic marker for a non-CNS disorder. Such markers are referred to herein as CNS "marker genes" or "disease surveillance genes" for non-CNS disease. It is understood, however, that the gene product of the marker gene can also serve as a diagnostic marker. In most cases, a plurality of differentially expressed markers are identified (e.g., a "profile" or "cluster" of markers is identified). The experimental animal is preferably an experimental mammal, and can be, e.g., an experimental rodent (e.g., a rat, mouse or guinea pig) or non-human primate (e.g., an ape, e.g., a monkey or chimpanzee).

The methods of detection of gene expression described herein, and particularly array and chip technology, are useful for methods of identifying CNS marker genes for non-CNS neoplasia. CNS samples are prepared from experimental and control animals (e.g., brains are biopsied or removed, or CSF samples are taken) and RNA, cDNA or protein is prepared from the samples as described herein. A single chip (e.g., a commercially available chip having probes for a large number of genes in the genome of the experimental animal species) can allow measurement of the level at which hundreds, thousands, or even tens of thousands of genes are expressed in the CNS sample of a test

experimental animal compared to a control experimental animal. Typically, clustering methodology or other bioinformatics tools are used to mine the data obtained from such large scale experiments and identify the genes or clusters of genes that are statistically significantly differentially expressed in an experimental sample compared to a control sample. Many such tools and programs are available to the skilled artisan. An exemplary method of data analysis is described herein and exemplified in the Examples below.

#### CNS Marker Genes for Neoplasia

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In one embodiment, identifying a CNS diagnostic marker for a non-CNS neoplastic disorder involves detecting changes in gene expression in the CNS in response to the presence of a non-CNS neoplasm in an experimental animal. For example, a neoplasm is induced in an experimental animal and gene expression in the CNS of the experimental animal is evaluated compared to a control animal. Methods for inducing growth of a non-CNS neoplasm, e.g., a cancer, in an experimental animal, are known in the art and include, e.g., chemical or radiation mutagenesis, or transplantation of a neoplastic cell (e.g., a neoplastic cultured cell or cell line) to the experimental animal. CNS genes or gene products whose expression is altered in the experimental animal compared to a control animal are identified as CNS markers or surveillance genes for neoplasia. Examples of CNS marker genes for cancer, particularly for carcinoma, are provided herein by FIGS. 2-26 and Examples 1-3.

# CNS Marker Genes for Rheumatoid Arthritis

In another embodiment, identifying a CNS diagnostic marker for rheumatoid arthritis (RA) involves detecting changes in gene expression in the CNS in an animal model of RA compared to a wild type animal. For example, the art-recognized rodent collagen induced arthritis (CIA) model can be used. In this model, arthritis is induced in a rodent, e.g., a DBA /1 mouse, by intradermal injection of purified collagen. 100 µg of purified type II collagen emulsified in complete adjuvant is typically injected at the base of the tail. Onset of arthritis is macroscopically visible as paw swelling or redness approximately three weeks after immunization (Williams et al., 1992, Proc. Natl. Acad. Sci. (USA), 89:9784-9788). Clinical features of arthritis are monitored by quantitatively

assessing paw swelling (e.g., with calipers) over a period of time. Severity of arthritis is assessed according to established clinical scores (Williams et al., 1995, Eur. J. Immunolo., 25:763-769). CNS genes or gene products whose expression is altered in the CIA animal compared to a control animal are identified as CNS markers or surveillance genes for RA.

Given the involvement of Th1 lymphocytes and B cells, proinflammatory cytokines, and a possible mimicry of bacterial LPS in disease evolvement, it is likely that genes that regulate these processes are candidates to be involved in early RA surveillance in the CNS. For example, proinflammatory cytokines produced in the brain such as IL-1β, TNF, IL-18, IFN-γ, IL-12, gp130; cytokines such as IL-6 and leukemia inhibitory 10 factor (LIF); neurotransmitters and neurotrophic factors such as N-methyl-D-aspartate (NMDA), brain-derived neurotrophic factor (BDNF), glial cell line-derived neurotrophic factor (GDNF), nerve growth factor (NGF); inhibitors of cytokines such as prostaglandin E2 (PGE2) and SOCS-1 and -3; SOCS regulators such as cAMP-inducing central peptides; brain molecules that are produced as a result of cytokine action, such as 15 pentraxin 3 (PTX3); hormone releasing factors such as cortocotropin; corticotropinreleasing hormone (CRH) and other hormones involved in the regulation of the HPA axis; pituitary corticotroph proteins such as POMC; molecules involved in NF-KBmediated signaling of inflammatory response; and other members of the families of these genes, as well as inducers and stimulators of these proteins, may be disease-surveillance 20 genes for RA. See, e.g., See, e.g., Blond et al., 2002, Brain Res., 958(1):89-99; Suk et al., 2001, Immunol. Lett., 77(2):79-85; Losy et al., 2001, Acta Neurol. Scand., 104(3):171-3; Opp et al., 2001, Neuroendocrinology, 73(4):272-84; Chesnokova et al., 2002, Endocrinology, 143(5):1571-4; Bousquet et al., 2002, Mol. Endocrinol., 15(11):1880-90; Polentarutti et al., 2000, J. Neuroimmunol., 106(1-2):87-94; Bayas et al., 25 2003, Neurosci. Lett. 335(3):155-8; Xu et al., 2000, Acta Pharmacol. Sin. 21(7):600-4; Fang et al., 2000, Neuroreport, 11(4):737-41).

# CNS Marker Genes for Asthma

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In another embodiment, identifying a CNS diagnostic marker for asthma involves detecting changes in gene expression in the CNS in an animal model of asthma compared

to a wild type animal. Several experimental models of asthma are known in the art, including rodent, sheep, and non-human primate models (for a review, see Isenberg-Feig et al., 2003, Curr. Allergy Asthma Rep. 3(1):70-8). Any of these can be used in the present methods. In one embodiment, the experimental model of asthma is performed according to Komai et al. (2003, Br. J. Pharmacol., 138(5):912-20). In brief, Balb/c mice are sensitized by intraperitoneal administration of 50  $\mu g$  of ovalbumin combined with 1 mg of alum (Al(OH)3) on day 0 and 12. From day 22 to 43 animals are exposed to daily aerosol challenges of 1% w/v of ovalbumin for 30 minutes. Control animals can include saline-injected animals and animals sensitized with ovalbumin and alum and challenged with saline. Airway function is evaluated by measuring one or more of: airway responsiveness to acetylcholine; IL-4, IL-5, and/or IL-13 levels; interferon- $\gamma$ levels; eosinophil numbers in bronchoalveolar fluids; specific IgG1 and IgG2a levels in sera; lung histology; and rectal temperature. CNS markers or surveillance genes for asthma are those whose expression is altered in the asthma model animal compared to a control animal, or those whose expression is altered after aerosol challenge compared to before aerosol challenge.

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Several gene products associated with the CNS have been shown to influence the Th-2 response and are candidates as disease-surveillance genes. These include glucocorticoid, one of the main hormonal mediators of stress, which acts on antigen-presenting cells to suppress the production of IL-12 *in vitro* and *ex vivo*; neurotransmitters norepinephrine or epinephrine; β-adrenoreceptor (ARs) agonists and antagonists (e.g., propranolol); modulators of neurotransmission such as adenosine and adenosine analogues; opiod system components, which influence the immunological response in general and the Th-1/Th-2 balance in particular; mediators of allergic reactions, such as histamine; neuropeptides such as substance P, vasoactive intestinal peptide and somatostatin, which increase the release of histamine from mast cells. See Blotta et al., 1997, J. Immunol. 158: 5589-5595; Elenkov et al., 1996, Proc. Assoc. Am. Physicians, 108: 374-381;Cooper et al., The biochemical basis of Neuropharmacology, Oxford University Press, 1996, p. 123; Link et al., 1999, J. Immunol. 164: 436-442; Loizzo et al., 2002, Br. J. Pharmacol., 135(5):1219-26; Lowman et al., 1988, British

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Journal of Pharmacology, Vol 95:121-130; and Elenkov et al., Annals of the New York Academy of Sciences, 2000, 917:94-105.

#### CNS Marker Genes for Diabetes

In another embodiment, identifying a CNS diagnostic marker for diabetes involves detecting changes in gene expression in the CNS in an animal model of diabetes compared to a wild type animal. Several experimental models of diabetes are known in the art, e.g., spontaneous models such as the NOD Mouse and BB Rat, and inducible models such as streptozotocin-induced (STZ) Diabetic Rats. These are reviewed in Cheta, 1998, J. Pediatr. Endocrinol. Metab., 11(1):11-9. CNS markers or surveillance genes for diabetes are those whose expression is identified to be altered in an induced animal compared to an uninduced animal (e.g., a streptozotocin-fed STZ rat compared to a control fed STZ rat), or those whose expression is altered in the early stages of spontaneous progression of disease.

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## CNS Marker Genes for Obesity

In yet another embodiment, identifying a CNS diagnostic marker for propensity for obesity involves detecting changes in gene expression in the CNS in an animal model of obesity, e.g., comparing CNS gene expression in an obesity-prone animal before and after obesity develops or is clinically detectable. The method can involve comparing differences in CNS gene expression between mouse strains that are either prone to obesity or resistant to obesity after being exposed to a fat-rich diet. For example, the method can employ the C57BL/KsJ(KsJ) or A/J strain of mice, both of which are resistant to the development of dietary obesity, or the obesity-prone strain C57BL/6J (B6).

Possible disease-surveillance genes for obesity or loss or body weight control include leptin, leptin receptor, ghrelin, cholecystokinin (CCK), CCK-A receptor, neuropeptide Y (NPY), proopiomelanocortin (POMC), α-melanocyte stimulating hormone (α-MSH), and other molecules that participate in the central control of energy balance. Given the fact that so many gene products orchestrate behaviors related to food intake, genetic deficiencies or the presence of particular polymorphic alleles in one or

more of these genes may induce disorders in the control of energy homeostasis leading to obesity. Such a deficiency or disruption in the normal signaling of such molecules can likely trigger an early signal that alters CNS gene expression.

# 5 <u>Isolating Homologous Sequences from Other Species</u>

The human homologs of CNS marker genes and their products (e.g., human homologs of CNS marker genes identified by experiments in non-human experimental animals) are useful for various embodiments of the methods described herein. Human homologs are known for most of the CNS marker genes provided herein. In those cases where a human homolog is not identified, several approaches can be used to identify such genes. These methods include low stringency hybridization screens of human libraries with a mouse marker gene nucleic acid sequence, polymerase chain reactions (PCR) of human DNA sequence primed with degenerate oligonucleotides derived from a mouse marker gene, two-hybrid screens, and database screens for homologous sequences.

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# Therapeutic Methods

The methods described herein can identify or diagnose the presence of a non-CNS disorder in a subject at an early stage in the pathogenic process. As such, the methods allow for early intervention, which can be the key to successful treatment and/or management of many disorders. For example, if a propensity for obesity or diabetes can be diagnosed at an early stage using the methods described herein, simple lifestyle or nutritional changes may be sufficient to stop or slow the progress of the disease, where such changes would not be sufficient if the disease were diagnosed at a later, more progressive stage. Similarly, a neoplasia that is detected at an early stage is more likely to be treated with less toxic therapeutic agents, or lower doses of a therapeutic agent, than would be used at a stage of advanced neoplasia, e.g., cancer.

#### Chemotherapeutic Agents

In one embodiment, the methods described herein can identify or diagnose the presence of a non-CNS neoplasia in a subject at an early stage, e.g., before a neoplasm has formed, before a neoplasm is clinically detectable, and/or before a tumor has become

malignant. As such, a neoplasm detected by a method described herein is amenable to treatment by an agent that targets neoplastic cells in general or targets specific neoplastic cells in particular. In one embodiment, a subject may be treated with a chemotherapeutic agent. Chemotherapeutic agents, as used herein, refer to chemical therapeutic agents or drugs used in the treatment of neoplasia. This term is used for simplicity notwithstanding the fact that other compounds may be technically described as chemotherapeutic agents in that they exert an anti-cancer effect. A number of exemplary chemotherapeutic agents are described below.

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Suitable chemotherapeutic agents include: antitubulin/antimicrotubule drugs, e.g., paclitaxel, taxol, tamoxifen, vincristine, vinblastine, vindesine, vinorelbin, taxotere; 10 topoisomerase I inhibitors, e.g., topotecan, camptothecin, doxorubicin, etoposide, mitoxantrone, daunorubicin, idarubicin, teniposide, amsacrine, epirubicin, merbarone, piroxantrone hydrochloride; antimetabolites, e.g., 5-fluorouracil (5-FU), methotrexate, 6-mercaptopurine, 6-thioguanine, fludarabine phosphate, cytarabine/Ara-C, trimetrexate, gemcitabine, acivicin, alanosine, pyrazofurin, N-Phosphoracetyl-L-Asparate=PALA, 15 pentostatin, 5-azacitidine, 5-Aza 2'-deoxycytidine, ara-A, cladribine, 5 - fluorouridine, FUDR, tiazofurin, N-[5-[N-(3,4-dihydro-2-methyl-4-oxoquinazolin-6-ylmethyl)-Nmethylamino]-2-thenoyl]-L-glutamic acid; alkylating agents, e.g., cisplatin, carboplatin, mitomycin C, BCNU=Carmustine, melphalan, thiotepa, busulfan, chlorambucil, plicamycin, dacarbazine, ifosfamide phosphate, cyclophosphamide, nitrogen mustard, 20 uracil mustard, and pipobroman, 4-ipomeanol; estrogen modulators, e.g., raloxifene; piroxicam; 9-cis retinoic acid.

Suitable dosages for the selected chemotherapeutic agent are known to those of skill in the art. For example, where the agent is doxorubicin, suitable dosage may include 30 mg/m<sup>2</sup> of patient skin surface area, administered intravenously, twice at 1 week intervals. However, one of skill in the art can readily adjust the route of administration, the number of doses received, the timing of the doses, and the dosage amount, as needed. Bearing in mind these considerations, generally, a suitable dose for a given chemotherapeutic agent is between 10 mg/m<sup>2</sup> to about 500 mg/m<sup>2</sup>, and more preferably, between 50 mg/m<sup>2</sup> to about 250 mg/m<sup>2</sup> of patient skin surface area (the skin surface of an average sized adult human is about 1.8 m<sup>2</sup>). Such a dose, which may be readily adjusted

depending upon the particular drug or agent selected, may be administered by any suitable route, including, e.g., intravenously, intradermally, by direct site injection, intraperitoneally, intranasally, or the like. Doses may be repeated as needed.

In one embodiment, because a method described herein can identify or diagnose the presence of a non-CNS neoplasia in a subject at an early stage, e.g., before a neoplasm has formed, before a neoplasm is clinically detectable, and/or before a tumor has become malignant, the dose of a chemotherapeutic agent may be lower than that typically used after a neoplasm, e.g., a cancer, is detected or diagnosed by clinical methods, such as visualization or palpation of a tumor mass.

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#### **Therapeutic Targets**

A CNS marker gene for a non-CNS disorder, e.g., a CNS marker gene described herein, may not only "sense" the presence of the disorder, but also actively participate in responding to the presence of the disorder by generating a response, e.g., an antitumor response. Alternatively, a CNS marker gene may respond to the presence of non-CNS disorder by promoting progression of the disorder, e.g., inducing growth of a neoplasm or promoting malignant transformation of a neoplasm. As a therapeutic strategy, one would want to promote the expression or activity of the former type of gene, and/or inhibit the expression of activity of the latter type of gene, in the CNS. Thus, regardless of whether a CNS marker gene generates a response to curb or promote a specific disorder, its identification can provide a target for inhibiting progression of the disorder.

One way to identify such CNS marker genes that are also potential therapeutic targets is to identify CNS genes that are differentially expressed in animals that exhibit an inhibitory response against a disease compared to animals that do not exhibit an inhibitory response. For example, experimental animals can be injected with tumor inducing cells (e.g., colon cancer cells such as CT26) that express an interleukin (IL), e.g., IL-12. Injection of tumor cells genetically modified to express IL-12 is known to induce Th1 immune mediated tumor rejection (Adris et al., 2000, Cancer Res., 60(23):6696-703). Control mice can be injected with tumor cells that do not express IL-12. At different times after injection, gene expression in the CNS is analyzed in the animals, as described herein, e.g., by microarray analysis. Thus, genes that "turn off" and

"turn on" specifically in the CNS (e.g., brain) of the animals can be identified. Some of these genes will respond to the presence of the IL. Others will correspond to genes actively engaged in the "stimulation" of the antitumor immune response. This strategy can be used for any interleukin gene that may be involved in the stimulation of an antitumor immune response. Identification of brain genes actively involved in "stimulating" an antitumor response will provide a target for therapeutic intervention, e.g., by direct use of the gene or its gene product, or by screening for agents that block or stimulate their activity.

A second strategy for identifying CNS genes that are potential therapeutic targets is by using transgenic animals (e.g., knockout mice) having brain specific disruptions (e.g., knockouts) in specific genes. A great number of CNS-specific knockout mice are currently available to the skilled artisan (see, e.g., the Jackson Laboratory web site, describing numerous JAX® mice models used in neurobiology), and many more can be expected to become routinely available. A role in the CNS response to non-CNS disease can be established for any particular gene for which a brain knockout animal can be obtained or produced, by inducing the disorder in the knockout mice (e.g., as described herein for cancer, RA, asthma or obesity), and evaluating disease outcome.

CNS marker genes and gene products that are also potential therapeutic targets are listed in FIG. 25A-E. These genes are or encode molecules involved in cell signaling, (e.g., growth factors, hormones, cytokines and their receptors) and are also differentially expressed markers in each of the tumors studied.

#### **Vaccines**

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The methods described herein also provide targets for preventive vaccination. A set of brain genes that "senses" a disease may include receptors for known or unknown ligands. A disease cell might produce these ligands to inhibit the induction of a brain-derived anti-disease response. In such an instance, identifying a CNS gene that is involved in an anti-disease response can lead to the identification of a gene product secreted by the diseased cell that might impact in the brain to inhibit disease response. A genetic vaccine targeting these products could be a viable therapeutic strategy.

One approach to identify CNS targets for preventive vaccination in the treatment of non-CNS disorders is the following: obtain a CNS gene expression profile (using techniques such as those described herein above) from animals that exhibit an anti-disease response, e.g., in the case of a tumor, an IL-12 mediated antitumor response, in an experimental tumor model. It is expected that from the cluster of genes "sensing" the tumor, some will change their expression levels in the presence of IL-12. This subset of genes will likely be those involved in "generating" the antitumor response. This subset of genes is likely to have predictable modulators. For example, if a CNS gene that changes its expression profile in response to a non-CNS gene in the presence of IL-12 is a receptor, one could predict that the change in gene expression of such a receptor could be brought about by its ligand. Thus, a preventive genetic vaccine could be designed to generate a memory response to such a ligand.

A second experimental approach can involve identifying those CNS genes that change their activity in response to a non-tumorigenic dose of tumor cells (e.g., a condition where neoplasia exists in the body, but no neoplasm is yet formed). From this subset of CNS genes one can predict the modulating genes responsible for their changes in activity, as explained above. Such modulating genes, which may be derived from the neoplastic cells, are likely to be initial tumor-derived signals of alarm in the peripheral body. Thus, a preventive genetic vaccine could be designed to generate a memory response to such genes.

A vaccine can be, e.g., a polypeptide or nucleic acid corresponding to the gene to be targeted. Vaccines described herein can be administered, or inoculated, to an individual in physiologically compatible solution such as water, saline, Tris-EDTA (TE) buffer, or in phosphate buffered saline (PBS). They can also be administered in the presence of substances (e.g., facilitating agents and adjuvants) that have the capability of promoting uptake or recruiting immune system cells to the site of inoculation. Vaccines have many modes and routes of administration. They can be administered intradermally (ID), intramuscularly (IM), and by either route, they can be administered by needle injection, gene gun, or needleless jet injection (e.g., Biojector<sup>TM</sup>, Bioject Inc., Portland, OR). Other modes of administration include oral, intravenous, intraperitoneal, intrapulmonary, intravitreal, and subcutaneous inoculation. Topical inoculation is also

possible, and can be referred to as mucosal vaccination. These include, for example, intranasal, ocular, oral, vaginal, or rectal topical routes. Delivery by these topical routes can be by nose drops, eye drops, inhalants, suppositories, or microspheres.

The following examples are illustrative only and not intended to be limiting.

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#### **EXAMPLES**

# Example 1: CNS Gene Expression Profiles Associated With Colon Carcinoma

CNS gene expression profiles associated with the presence of a peripheral tumor were identified using gene expression microarray analysis on brain tissue from experimental animals implanted peripherally with tumor cells. This example describes the identification of brain gene expression profiles associated with colon carcinoma.

Male BALB-C mice were injected subcutaneously with 5 x 10<sup>5</sup> CT-26 WT cells, a murine colon carcinoma cell line (ATCC cat # : CRL-2638), resuspended in 300 μl of PBS, as described below. Control mice were injected with the corresponding volume of PBS following the same procedure. After a specified time, the animals were sacrificed, their brains dissected, and first strand cDNA was synthesized from total polyA+RNA prepared from different brain regions, as described in detail below. Gene expression microarray analysis was performed with the first strand cDNA by hybridizing to preprinted slides (Corning's CMT-GAP™ II Coated Slides) containing Pan® Mouse 10K Oligo set A (MWG Biotech). This slide set contains probes for 10,000 genes selected from mouse genes that have been functionally defined.

The data from the microarray experiments was analyzed with a Virtek® ChipReader® laser scanner model A0-B0-05 (Virtek Vision Corp, Waterloo, ON, Canada) using the Virtek ChipReader v2.0 software, as described in more detail below.

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# **Experimental Methodology**

Cell Lines: The experimental work was based on the following murine cell lines: CT26WT colon carcinoma (ATCC cat #: CRL-2638), LL/2(LLC1) lung carcinoma (ATCC cat #: CRL-1642) and 4T1 breast carcinoma (ATCC cat #: CRL-2539). All cell lines were grown in P-100 plates with 10 ml of the corresponding medium. All culture media were sterilized by filtration using 0.22  $\mu$ m CA filter. CT-26 cells were grown in

DMEM containing 1.5 g/L Sodium Bicarbonate, 10 mM Hepes, and 1 mM Sodium pyruvate, supplemented with 10% Fetal Bovine Serum at 37°C with 5% CO<sub>2</sub>. LL/2(LLC1) cells were grown in DMEM containing 4.5 g/L Glucose, 1.5 g/L Sodium Bicarbonate, 10 mM Hepes, and 1 mM Sodium pyruvate, supplemented with 10% Fetal Bovine Serum at 37°C with 5% CO<sub>2</sub>. 4T1 cells were grown in RPMI 1640 containing 4.5 g/L Glucose, 1.5 g/L Sodium Bicarbonate, 10 mM Hepes, and 1 mM Sodium pyruvate, supplemented with 10% Fetal Bovine Serum at 37°C with 5% CO<sub>2</sub>.

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In vivo studies: Six week-old animals were housed in an Hepa filtered air rack, 5 animals per cage (both tumor and control animals in the same cage) with food and water ad libitum. Balb-C males were injected subcutaneously with 5 x  $10^5$  CT-26 WT cells resuspended in 300  $\mu$ l of PBS. BALB-C female mice were injected subcutaneously with 1 x  $10^5$  4T-1 cells resuspended in 100  $\mu$ l of PBS. C-57/BL6 male were injected subcutaneously with  $1 \times 10^6$  LL/2(LLC1) cells resuspended in 300  $\mu$ l of PBS. Control animals were injected with the corresponding volume of PBS following the same procedure.

For each tumor type 2 different experiments were performed and 3 time points evaluated in duplicate. Each single time point corresponded to 15 mice. All injections were done using a 27-G syringe. At the corresponding time, mice were killed by cervical dislocation. Mice were immediately decapitated, the brain extracted and dissected using the following procedure: the hypothalamus and the cerebellum were dissected, the brain was cut with a surgical razor blade leaving the right and left hemispheres separated, and two persons dissected the midbrain, the hippocampus, the prefrontal cortex and the striatum from each brain hemisphere. All brain regions were immediately frozen in dry ice and stored at  $-80^{\circ}$ C until RNA extraction.

<u>Preparation of Poly A+ RNA</u>: Poly A+ RNA was obtained from total RNA using the MicroPoly(A) Pure® kit from Ambion. In general, starting material was 400 μg total RNA to which a volume of 5M NaCl was added up to a final concentration of 0.45 M NaCl. After mixing, samples were transferred to an RNase-free microfuge tube. After adding binding buffer provided by the manufacturer, the RNA was heated for 5 minutes at 65°C and immediately chilled on ice for 1 minute. Oligo (dT) Cellulose was added to the sample, mixed by inversion and incubated for 60 minutes at room temperature with

gentle agitation. This was followed by centrifugation at 4,000 rcf for 3 minutes. After the supernatant was removed, the pellet was treated with 1 ml binding buffer, mixed and spun down by centrifuging at 4,000 rcf for 3 minutes. After removing the supernatant, the pellet was washed 3 times with binding buffer followed by 4 washes with wash buffer. The Oligo(dT) Cellulose was then dissolved in 400 µl of wash buffer provided by the manufacturer and transferred to a spin column when the resin was washed 4 more times. When the flow-through of the column reached an absorbance of <0.05 OD at A260, the mRNA was eluted from the Oligo(dT) Cellulose with 200 µl of Elution Buffer (provided by the manufacturer) pre-warmed at 65°C. The eluted polyA+ RNA was concentrated with a mixture containing 20 µl of 5 M Ammonium Acetate, 1 µl Glycogen and 550 µl of 100% ethanol. After overnight precipitation at -20°C samples were centrifuged at 14,000 rcf for 20 minutes at 4°C. After careful removal of the supernatant the pellet containing the polyA+ RNA was resuspended in 10 µl of DEPC treated Water/EDTA.

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Labeling of probes for microarray hybridization: Labeling was performed by two indirect methods. The first method used aminoallyl labeled nucleotides via first strand cDNA synthesis using SuperScript Reverse Transcriptase followed by coupling of the aminoallyl to either Cyanine 3 or 5 (Cy3/Cy5) fluorescent molecules (Amersham Pharmacia). To 3 µg of poly(A+) RNA were added 0.6 µl Random Primers (pd (N)6, Invitrogen) (3  $\mu$ g/ $\mu$ l) and 1.2  $\mu$ l Oligo (dT)12-18 (0.5  $\mu$ g/ $\mu$ l). Milli-Q H<sub>2</sub>O was added up to a final volume of 15.5  $\mu$ l. The mixture was heated at 65°C for 5 min, chilled on ice and spun down. 12.5 µl of a Master Mix containing: 6 µl of 5X First Strand Buffer, 3 µl of 100 mM DTT, 0.6 µl of 50X aminoallyl (Sigma Co)-dNTP mix (Amersham Pharmacia), 1.5 μl of Rnase OUT (40 units/μl, Invitrogen), 1.4 μl Milli-Q H<sub>2</sub>O were added to each tube, incubated at 37°C for 2 minutes, followed by the addition of 2  $\mu l$  of SuperScript II RT (Invitrogen). After incubation for 2 hr at 37°C tubes were transferred 15 min at 70°C. At the end, tubes were spun down. RNA was degraded by the sequential addition of 3  $\mu$ l 2.5 M NaOH incubated at 37°C for 15 min, then 15  $\mu$ l of 2 M HEPES free acid, 4.8 µl 3 M NaAcO (pH 5.2) and finally 150 µl of 100% EtOH. After mixing, tubes were incubated at -20°C for 1 hr. Tubes were centrifuged for 30 min at 4°C, the supernatant was removed and the pellet was washed twice in 70% ethanol. The pellet

was dissolved in 2.25 µl Milli-Q H<sub>2</sub>O. Coupling of fluorescent Cy3 and Cy5 was performed by initially adding 2.25 µl of 0.2 M NaHCO3 (pH 9.0) and then 4.5 µl of the DMSO/dye mixture to the 4.5 µl cDNA sample. Tubes were mixed well and incubated for 1 hr at room temperature in the dark. For probe purification 500 µl of Loading Buffer were added to the sample and mixed. A SNAP Column (Invitrogen) was placed on a collection tube and the sample loaded on the column and incubated at room temperature for 2-5 min. The system was centrifuged at maximum speed for 1 min and the flow-through was discarded. After two more washes the SNAP column was put back in the collection tube and centrifuged at maximum speed for 30 sec to remove residual Wash Buffer from the membrane filter. cDNA was eluted by adding 60 µl TE buffer to the SNAP column, incubated for 2-5 and centrifuged at maximum speed at room temperature for 1-2 min. After saving the first eluate, the elution was repeated and both samples were combined.

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Alternatively, labeling of poly A+ RNA was performed with a Clontech kit following manufacturer instructions. Briefly, to 3  $\mu g$  of poly(A+) RNA were added 15 0.6  $\mu$ l Random Primers (pd(N)6) (3  $\mu$ g/ $\mu$ l), 0.5  $\mu$ l Oligo(dT)12-18 (0.5  $\mu$ g/ $\mu$ l), and deionized H<sub>2</sub>O up to 25 µl. After heating at 70°C for 5 min the tubes were placed at 37°C, and then 25  $\mu$ l of Master Mix were added (10  $\mu$ l of 5X cDNA Synthesis Buffer,  $5~\mu l$  of 10X dNTP Mix, 7.5  $\mu l~H_2O$  and 2.5  $\mu l~MMLV$  Reverse Transcriptase (200 units/µl). Tubes were incubated at 37°C for 1 hr, followed by 5 min at 70°C. After 20 few minutes at 37°C, RNA was eliminated by adding 0.5 μl RNase H for 15 min at 37°C and then 0.5 µl of 0.5 M EDTA (pH 8.0) together with 5 µl of QuickClean resin. After inserting a 0.45-µm Spin Filter into the collection tube, the sample was transferred into the Spin Filter. The cDNA was concentrated with 3M Sodium Acetate, the addition of ice-cold 100% ethanol and centrifugation at maximal speed for 20 min at 4°C. The pellet 25 was washed once in 70% ethanol, air dried and dissolved in 10 µl 2X Fluorescent Labeling Buffer. Fluorescent dye coupling was performed by adding 10 µl of the DMSO/dye mixture to  $10 \mu l$  of the cDNA sample. This mixture was mixed well and incubated for 30 min in the dark. 2 µl of 3M Sodium Acetate and 50 µl of 100% ethanol were added. After 2 hr at -20°C the tube was centrifuged for 20 min. After washing the 30

pellet once in 70% ethanol the pellet was dissolved in 100  $\mu$ l  $H_2O$ . Probes were purified on NucleoSpin columns.

Quantification of the levels of incorporation of dyes and total DNA: The extent of dye incorporated was obtained by the absorbance at 550 nm and 650 nm for Cy3- and Cy5-probes, respectively. The amount of DNA was obtained by the absorbance at 260 nm. At the end of the entire procedure the amount of total DNA obtained was 0.34-0.65  $\mu$ g DNA / 1  $\mu$ g poly A+ RNA for the Clontech procedure and 0.8 – 1.2  $\mu$ g DNA / 1  $\mu$ g poly A+ RNA for the SuperScript and indirect labeling procedure. The current percentage of dye incorporation was 5 – 15 % in the first case and 7.5 – 20 % in the second.

# Microarrays and Data Analysis

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Prehybridization: The Prehybridization Buffer (5 ml of 20X SSC Buffer, 0.25 ml of 20% SDS, 5 ml of 10% BSA and 24.75 ml of Milli-Q H<sub>2</sub>O) was preheated at 42°C. The printed slide was put in a 50 ml-Falcon polypropylene tube containing the preheated prehybridization buffer and incubated at 42°C for 40 min. After washing five times, 1 min each, with Milli-Q H<sub>2</sub>O preheated at 42°C in a Wash Station, slides were washed four or five times in 2-propanol. The slide was dried by centrifugation for 1 min using a Microarray Centrifuge. Cover glasses were washed with Milli-Q H<sub>2</sub>O and 2-propanol and dried. Slides were used immediately for hybridization.

<u>Probe preparation</u>: 2 μg of probe (1 μg Cy3- + 1 μg Cy5-labeled probes) were used per slide. This amount represents 100-110 pmoles and 70-80 pmoles of Cy3 and Cy5 incorporated dye, respectively. If dye incorporation levels were below that value, the amount of nucleotide was increased to reach these values (picomoles of labeled probe). Probe was concentrated by speedvac to about 20  $\mu$ l, combined and mixed well.

Hybridization: Each hybridization mix contained: 20 μl of 4 X Hybridization Buffer (Amersham Pharmacia; Cat. No. RPK0325), 24 μl of formamide (final concentration, 30 %) and 16 μl of Salmon Sperm DNA (final concentration 1μg / μl). This master mix was added to probes, mixed well, heated at 95°C for 3 min, snap cooled on ice for 1 min and centrifuged at 16.000 x g for 1 min. A pre-hybridized microarray slide (array side up) was placed in a hybridization chamber. Two Parafilm strips were put

at both sides of the array printed area. Finally, the probe was placed carefully on the top of the slide surface followed by the coverslip on top of it (FIG. 1).  $10 \,\mu l$  of Milli-Q  $H_2O$  (20  $\mu l$  total) was added to the small wells at each end of the chamber to seal the chamber. Slides were incubated at 42°C for 16-20 hours under gently mixing in a 3D-rotator. At the end of the hybridization, the slide was carefully removed and washed with washing buffer preheated at 42°C for 5 min with agitation (2 X SSC, 0.1 % SDS). Slides were washed twice more in different chambers, each time for 5 minutes (First in 1 X SSC and then in 0.1 X SSC). The slide was dried by centrifugation for 1 min in a microarray centrifuge and placed in a light tight slide box until scanning.

Data acquisition and image processing: The slides were scanned with a Virtek ChipReader laser scanner model A0-B0-05 (Virtek Vision Corp, Waterloo, ON, Canada) using the Virtek ChipReader v2.0 software. Three images were obtained for each of the Cy3 and Cy5 channels with different detector sensitivity values for each slide, with a resolution of 10 μm and a pixel depth of 16 bits. The images were stored as 16 bit TIFF files (Tagged Image File Format) and analyzed with Virtek ChipReader v2.0 software. The results were stored in plain text files with the following fields separated by tabulations: GridName, Column#, Row#, CentroidX, CentroidY, SNR, Signal Average, Signal Median, Signal Std, Signal pixels, Background Average, Background Median, Background Std.

Data filtration and normalization: All the data processing was performed under the R System v1.6.2 (R Development core Team<sup>TM</sup> software). The data was filtered to eliminate background data points (spots with size less than 75 pixels or with a mean to median correlation less than 80% (Tran and Peiffer, 2002, Nucleic Acids Res. 30(12), e54), to eliminate saturated data points (spots with a proportion of saturated pixels greater than 20%), and to eliminate low signal data points (spots with signal to noise ratio below 2). The signal was corrected for background and the signal volume was estimated as (Signal Average – Background Average) x Signal pixels. The base 2 logarithm of the ratio and the product between Cy5 and Cy3 was calculated as:

$$M = \log_2(\text{Cy5/Cy3})$$
 (1)

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$$A = \log_2(\text{Cy5} \times \text{Cy3}) \quad (2)$$

Data was normalized using locally weighted linear regression of M vs. A over rank consistency filtered data for each print tip (rank consistent print tip lowest fit) for 150 or more data points, or using a rank consistent global lowest fit algorithm when the number of data points was below 150.

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For the rank consistency filter, the rank of Cy3 and Cy5 intensities of each gene on the slide were separately calculated. For a given gene, if the ranks of Cy3 and Cy5 intensities differed by less of a threshold value d, this gene was classified as rank consistent. This process was iteratively repeated until the number of rank consistent genes did not change. The threshold level d was defined as follows:

$$d = p \cdot (1 + 1/i) \cdot fn(A) \cdot n \tag{3}$$

where i is the iteration number; n is the number of data points, which is equal to the number of spots in the slide for the first iteration cycle, and equal to the number of rank consistent genes defined in cycle i-1 for the next cycles; fn(A) is the normal density function with mean equal to the average of A and standard deviation (SD) equal to the estimated SD of A; and p is a proportionality constant that was set to 0.5.

Then, the value of M for each value of A that follows the central tendency of the data (Mc) was estimated from the rank consistency data with the R package lowest function, and it was subtracted from the empirical M value to obtain the normalized M data (M'). From this point, all further analysis was performed with the normalized M' data.

Outlier data points were eliminated from the triplicate data with a leave-one-out algorithm. Briefly, a data point was discarded as being outlier if it was outside the confidence interval defined by the other two data points with a confidence level of 95%.

A gene expression data set was generated for each slide with the average of nonoutlier data points.

Differences in scale between Cy3 and Cy5 channels can lead to an asymmetric distribution of M' data. To correct this deviation, M' data was transformed to be normally

distributed. First, a uniform distributed data set between 0 and 1 was obtained with the transformation Mu = rank(M')/(n+1), where n is the number of data points. Then, a normal distributed data set, with mean equal to the median of M' and SD equal to the estimated SD of M', was obtained with the transformation: Mn = qnorm(Mu), where qnorm is the normal quantile function included in the R package.

Data integration between replicated slides: Each labeled probe was hybridized at least twice. If the scale (i.e. variance) between replicated slides was different (p < 0.05, Fligner-Killeen test for homogeneity of variances), data were transformed to be equally scaled. Assuming that the ratios follow a normal distribution with mean zero and variance  $a_i 2\sigma 2$ , we estimated  $a_i$  as follows:

$$a_i = \frac{MAD_i}{\sqrt[l]{\prod_{i=1}^{I} MAD_i}}$$
 (4)

with I denoting the total number of slides, and the median absolute deviation MAD defined by,

$$MAD = \text{median}_j | M_{ij} - \text{median}_j (M_{ij})$$
 (5)

where  $M_{ij}$  denotes the  $j^{th}$  spot in the  $i^{th}$  slide.

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Outlier data points were eliminated from three or more replicated data sets with a leave-one-out algorithm as described above, and an integrated data set was obtained with the averaged M values from non-outlier data.

Analysis and integration of replicated experimental data and noise analysis: Experiments were performed at least twice. If the scale (i.e. variance) between replicated experiments was different (p < 0.05, Fligner-Killeen test for homogeneity of variances), data were transformed to be equally scaled.

Outlier data points were eliminated from three or more replicated data sets with a leave-one-out algorithm. The arithmetic mean (Mn) and SD were estimated from the non-outlier data set. A noise sampling method was used for p-value estimation (Draghici et al., Noise sampling method: an ANOVA approach allowing robust selection of

differentially regulated genes measured by DNA microarrays, Bioinformatics, in press). Briefly, an estimation of the noise is obtained from the replicated data as the difference between the ratio expression for gene g in experiment e and the mean for gene g among experiments. Because noise varies with intensity: low intensity spots tend to have more noise than high intensity ones, the intensity range was divided in bins, and noise distributions constructed for each such bin. Assuming that the noise distribution is normal, which was the case for most experiments, it was mapped from the distribution of the noise to the distribution of the log ratios by the scaling factor 1 / sqrt (n - 0.5), where n is the number of replicates.

Cluster analysis: Before cluster analysis, the data was scaled as follows:

Ms = (M - Mn(M)) / SD(M). A figure of merit algorithm (Yeung et al., 2001,

Bioinformatics, 17(4):309-18) was used to identify the clustering algorithm and the

number of clusters that minimized the intra-cluster variability. After examining the figure

of merit of all the datasets analyzed with seven different clustering algorithms and

different variations of such algorithms that led to a total of 51 different clustering

methods, it was decided to subset the data in 12 clusters with a hierarchical algorithm

using euclidean distance between gene expression patterns and a Ward's minimum

variance agglomeration method (Hartigan (1975). Clustering Algorithms. New York:

Wiley.). Genes with similar expression patterns among the experiments were clustered

together using routine hierarchical clustering techniques.

#### Results

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After quality filtering and normalizing the microarray data, sequences with a p-value below 0.05 were identified as differentially expressed (DE). Further analysis was performed on this subset of sequences to select and cluster sequences according to specific criteria. Genes with similar expression patterns among the experiments were clustered together using hierarchical clustering techniques as described above.

Cluster analysis I: A first clustering analysis (cluster analysis I) identified DE sequences (p < 0.05) up- or down-regulated in one of two experimental time points tested (72 and 192 hours). These are shown for the colon cancer model in FIG. 2 (DE sequences in midbrain), FIG. 8 (DE sequences in cortex), FIG. 14 (DE sequences in striatum), and FIG. 18 (DE sequences in hypothalamus). Cluster graphs (the last sheet of

each Figure) show whether genes in a particular cluster were up- or down-regulated (y-axis) at each time point tested (x-axis). For example, in FIG. 2-8, clusters 4-6 were up-regulated at 72 hours (x-axis 1.0) and clusters 9-12 were up-regulated at 192 hours (x-axis 2.0). In FIG. 8-13, clusters 9 and 10 were down-regulated at 192 hours while clusters 11 and 12 were up-regulated at 192 hours. In FIG. 14-8, clusters 4, 5 and 6 are close to midline at 72 hours and down-regulated at 192 hours. FIG. 18-8 shows clusters 11 and 12 up-regulated at 192 hours.

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Cluster analysis II: A more stringent clustering analysis (cluster analysis II) revealed DE sequences (p < 0.05) up- or down-regulated in both experimental time points tested. These are shown in FIG. 3 (DE sequences in midbrain), FIG. 9 (DE sequences in cortex), FIG. 15 (DE sequences in striatum), and FIG. 19 (DE sequences in the hypothalamus). For example, FIG. 3-2 shows that only cluster 7 is up-regulated at both time points, while the rest of the clusters are down-regulated at both time points. FIG. 19-2 shows that only cluster 4 is up-regulated at both time points.

<u>Secreted markers</u>: In a third analysis, the filtered data were reclustered to select sequences that should correspond to a secreted product and have a p value for differential expression below 0.05 (p<0.05). The results of this analysis for colon cancer is shown in FIG. 24(A). FIG. 24(A) lists markers corresponding to secreted products, that were differentially expressed in the colon cancer model at any time point studied. Secreted markers are particularly useful in that their expression can be detected in cerebral or cerebrospinal fluid, avoiding the need for a solid tissue biopsy.

Shared DE markers: In a final analysis, the filtered data were reclustered to select sequences that were differentially expressed in all tumors analyzed. Seven DE sequences were found to be shared among all three carcinomas studied. These were hepatocyte growth factor (HGF), apherin A3, chemokine (C-C motif) ligand 4, growth differentiation factor-9b (GDF-9b); bone morphogenetic protein 15 (BMP 15), neuroblastoma suppressor of tumorigenicity 1, melanocyte proliferating gene 1, and fibroblast growth factor 22 (FGF 22). (FIG. 26)

# Example 2: CNS Gene Expression Profile Associated With Breast Carcinoma

This example describes the identification of brain gene expression profiles associated with breast carcinoma.

BALB-C mice were injected subcutaneously with 1 x  $10^5$  4T-1 breast carcinoma cells (ATCC cat #: CRL-2539) resuspended in 100  $\mu$ l of PBS. All experimental methods, microarrays and data analysis were otherwise performed as described above for Example 1.

#### Results

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Quality filtering, normalization and analysis of the microarray data was performed as discussed above.

Cluster analysis I: a first clustering analysis (cluster analysis I) identified DE sequences (p < 0.05) up- or down-regulated in only one of the three experimental time points tested (18, 72 and 192 hours). These are shown for the breast cancer model in FIG. 4 (DE sequences in midbrain), FIG. 10 (DE sequences in cortex), and FIG. 20 (DE sequences in hypothalamus). For example, FIG. 4-16 shows that the genes of cluster 1, 2 and 3 were up-regulated at 72 hours (x-axis 2.0). In FIG. 10-13, clusters 2 and 3 show up-regulation at 72 hours. In Fig. 20-21, only clusters 5-7 are down-regulated at 192 hours (x-axis 3.0).

Cluster analysis II: a more stringent clustering analysis (cluster analysis II) revealed DE sequences (p < 0.05) up- or down-regulated in at least two of the three experimental time points. These are shown in FIG. 5 (DE sequences in midbrain), FIG. 11 (DE sequences in cortex), and FIG. 21 (DE sequences in hypothalamus). In FIG. 5, only cluster 7 shows up-regulation at any time point, while the remaining clusters are generally down-regulated. Similarly, only one cluster (cluster 6) is up-regulated in FIG. 11-6. Only clusters 10-12 are up-regulated in FIG. 21-7.

<u>Secreted markers</u>: in a third analysis, the filtered data were reclustered to select sequences that correspond to a secreted product and have a p value for differential expression below 0.05 (p<0.05). The results of this analysis for breast cancer is shown in FIG. 24B. FIG. 24B lists markers corresponding to secreted products, that were differentially expressed in the breast cancer model at any time point studied.

# Example 3: CNS Gene Expression Profile Associated With Lung Carcinoma

This example describes the identification of brain gene expression profiles associated with lung carcinoma.

Male C-57/BL6 mice were injected subcutaneously with 1x10<sup>6</sup> lung carcinoma LL/2(LLC1) cells (ATCC cat #: CRL-1642) resuspended in 300 µl of PBS. All experimental methods, microarrays and data analysis were otherwise performed as described above for Example 1.

#### Results

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Quality filtering, normalization and analysis of the microarray data was performed as discussed above.

Cluster analysis I: a first clustering analysis (cluster analysis I) identified DE sequences (p < 0.05) up- or down-regulated in only one of the three experimental time points tested (18, 72 and 192 hours). These are shown for the lung cancer model in FIG. 6 (DE sequences in midbrain), FIG. 12 (DE sequences in cortex), FIG. 16 (DE sequences in striatum), and FIG. 22 (DE sequences in hypothalamus). For example, FIG. 6-15 shows that clusters 2, 3 and 4 are up-regulated at 72 hours (x-axis 2.0) while clusters 5-11 are up-regulated at 18 hours (x-axis 1.0). FIG. 12-13 shows that clusters 2, 3 and 4 are up-regulated at 72 hours. In FIG. 22-21, only clusters 3, 4, 8 and 9 are down-regulated at 192 hours.

Cluster analysis II: a more stringent clustering analysis (cluster analysis II) revealed DE sequences (p < 0.05) up- or down-regulated in at least two of the three experimental time points. These are shown in FIG. 7 (DE sequences in midbrain), FIG. 13 (DE sequences in cortex), FIG. 17 (DE sequences in striatum), and FIG. 23 (DE sequences in hypothalamus). In FIG. 7-5 and FIG. 17-5, all the clusters, except for cluster 12 in each set, are down-regulated at every time point studied. FIG. 13-6 shows that all but two clusters (11 and 12) were down-regulated.

<u>Secreted markers</u>: in a third analysis, the filtered data were reclustered to select sequences that correspond to a secreted product and have a p value for differential expression below 0.05 (p<0.05). The results of this analysis for lung cancer is shown in FIG. 24C. FIG. 24C lists markers corresponding to secreted products, that were differentially expressed in the lung cancer model at any time point studied.

# Example 4: Diagnosis of Cancer in a Human by Detecting a Gene Product Profile

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This example describes a diagnostic test for non-CNS carcinoma performed on a human subject. The subject is a carrier of the BRCA1 breast cancer susceptibility gene.

A CSF sample is obtained from the subject by means of a lumbar puncture. This procedure is done on an outpatient basis under local anesthetic. The CSF sample is used immediately in the diagnostic assay, or is cooled or frozen and stored or transported to a facility where the diagnostic test is performed.

The diagnostic test involves contacting the CSF sample to an antibody array containing a panel of 25 antibodies that can detect a set (cluster) of CNS gene products that are associated with the presence of breast cancer when secreted in a characteristic profile in the CSF. The panel includes antibody probes for one or more CNS markers for breast carcinoma listed in FIG. 24B. Thus, in this example, the characteristic profile is the CNS "reference profile" for breast carcinoma.

The results of the antibody array are obtained by routine techniques, such as fluorescence detection and measurement of bound antibody vs. unbound antibody for each position (each antibody) on the array. A dataset of the value for the level of each polypeptide detected in the CSF sample by each antibody on the array is generated. The dataset is used directly as the test expression profile or the dataset is converted into a format comparable to the format of the reference profile to which the test profile is compared.

Once the test expression profile is generated, the test profile is compared to the reference expression profile. In this example, the reference profile is a dataset that includes relative values of expression for a panel of 10 CNS gene products secreted into the CSF, all of which are known to be down-regulated at least 30%, on average, in subjects who have early stage breast cancer. The gene products include one or more of the gene products shown in FIG. 4, 10, 20, 5, 11, 21 or 24B. If the test profile shows that 7 or more of the genes in the panel are down-regulated by at least 20% in the test sample, the test profile matches the reference profile and the subject is determined to have (or be at risk for) early stage breast cancer.

#### OTHER EMBODIMENTS

A number of embodiments of the invention have been described. Nevertheless, it will be understood that various modifications may be made without departing from the spirit and scope of the invention. Accordingly, other embodiments are within the scope of the following claims.

# We claim:

- 1. A method of diagnosing a non-central nervous system (non-CNS) disorder in a subject, the method comprising:
- detecting expression of a gene in a CNS sample of the subject, and correlating the result of the detecting step to the presence or absence of a non-CNS disorder.
- 2. The method of claim 1, further comprising the step of obtaining the CNS sample.
  - 3. The method of claim 1, wherein the CNS sample is one or more brain cells.
  - 4. The method of claim 3, wherein the brain cells are selected from the group consisting of cells from: the hypothalamus, the midbrain, the prefrontal cortex and the striatum.
    - 5. The method of claim 1, wherein the CNS sample is cerebrospinal fluid.
- 6. The method of claim 1, wherein the non-CNS disorder is selected from the group consisting of: cancer, rheumatoid arthritis, asthma, diabetes and obesity.
  - 7. The method of claim 1, wherein the non-CNS disorder is a carcinoma.
- 8. The method of claim 1, wherein the non-CNS disorder is a solid tumor less than 0.5 cm in diameter.
  - 9. The method of claim 1, wherein the gene encodes a gene product selected from the group consisting of: a hormone, a growth factor, an immune system component, a cytokine.

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- 10. The method of claim 7, wherein the gene encodes a gene product listed in any of FIGS. 2-26, or a human or other mammalian homolog thereof.
- 11. The method of claim 7, wherein the gene encodes a gene product selected from
  5 the group consisting of: hepatocyte growth factor (HGF), apherin A3, chemokine (C-C
  motif) ligand 4, growth differentiation factor-9b (GDF-9b); bone morphogenetic protein
  15 (BMP 15), neuroblastoma suppressor of tumorigenicity 1, melanocyte proliferating
  gene 1, and fibroblast growth factor 22 (FGF 22).
- 10 12. The method of claim 1, wherein detecting expression of the gene comprises detecting the mRNA corresponding to the gene.
  - 13. The method of claim 1, wherein detecting expression of the gene comprises detecting a polypeptide product encoded by the gene.

14. The method of claim 1, wherein detecting comprises detecting expression of a plurality of genes in a CNS sample of the subject

- 15a. The method of any one of the preceding claims, wherein the detecting step comprises performing a microarray assay.
  - 15b. The method of claim 1, wherein the subject is a human.

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16. A method of diagnosing a non-central nervous system (non-CNS) disorder in a subject, the method comprising:

obtaining a test gene expression profile for two or more CNS genes from the subject; and

comparing the test gene expression profile with a reference gene expression profile associated with the presence of a non-CNS disorder, wherein a test gene expression profile that matches the reference gene expression profile indicates the subject has a non-CNS disorder.

17. The method of claim 16, further comprising generating a record of the result of the comparing step; and optionally transmitting the record to the subject, health care provider or other party.

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- 18. The method of claim 16, wherein the non-CNS disorder is selected from the group consisting of: cancer, rheumatoid arthritis, asthma, diabetes and obesity.
- 19. The method of claim 16, wherein the non-CNS disorder is a carcinoma.

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- 20. The method of claim 16, wherein the non-CNS disorder is a solid tumor less than 0.5 cm in diameter.
- 21. The method of claim 16, wherein at least one of the two or more CNS genes is selected from the group consisting of: a hormone, a growth factor, an immune system component, and a cytokine.
  - 22. The method of claim 19, wherein at least one of the two or more CNS genes encodes a gene product listed in FIGS. 2-26, or a human or other mammalian homolog thereof.
  - 23. The method of claim 19, wherein at least one of the two or more CNS genes encodes a gene product selected from the group consisting of: hepatocyte growth factor (HGF), apherin A3, chemokine (C-C motif) ligand 4, growth differentiation factor-9b (GDF-9b); bone morphogenetic protein 15 (BMP 15), neuroblastoma suppressor of tumorigenicity 1, melanocyte proliferating gene 1, and fibroblast growth factor 22 (FGF 22).
- The method of claim 16, wherein the step of obtaining the test gene expression
   profile comprises detecting mRNA corresponding to the two or more CNS genes.

- 25. The method of claim 16, wherein the step of obtaining the test gene expression profile comprises detecting polypeptide products encoded by the two or more CNS genes.
- The method of claim 16, comprising obtaining a test gene expression profile for a
   plurality of CNS genes.
  - 27. The method of any one of claims 16-26, wherein the step of obtaining the test gene expression profile comprises performing a microarray assay.
- A method of treating a subject, the method comprising: diagnosing a non-central nervous system (non-CNS) disorder according to the method of claim 1 or 16; and administering to the subject a therapeutic agent for the disorder.
- 15 29. The method of claim 28, wherein the therapeutic agent is chemotherapeutic agent.
  - 30. The method of claim 29, wherein the chemotherapeutic agent is selected from the group consisting of: an antitubulin/antimicrotubule drug, a topoisomerase I inhibitor, an antimetabolite, and an alkylating agent.

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31. A method of identifying a diagnostic marker for a non-central nervous system (non-CNS) disorder in a human, the method comprising:

inducing a non-CNS disorder in a test experimental animal;

comparing expression of a gene in a CNS sample from the test experimental animal to expression of the gene in a CNS sample from a control experimental animal; and

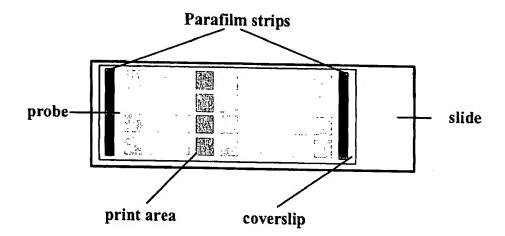
selecting as a diagnostic marker a human homolog of a gene that is differentially expressed in the CNS sample from the test experimental animal compared to the CNS sample from the control experimental animal.

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- 32. The method of claim 31, wherein a non-CNS neoplasm is induced by chemical or radiation mutagenesis.
- 33. The method of claim 31, wherein a non-CNS neoplasm is induced by
  administering a neoplastic cell to the experimental animal.
  - 34. The method of claim 31, wherein the experimental animal is an animal model of rheumatoid arthritis, diabetes, asthma, obesity or diabetes.
- 10 35. The method of claim 31, wherein the experimental animal is a non-human primate.
  - 36. The method of claim 1, 16 or 28, wherein the subject lacks a clinical sign of a disorder as evaluated by imaging analysis.
  - 37. The method of claim 1, 16 or 28, wherein the subject has a family history of the disorder.
- 38. The method of claim 1, 16 or 28, wherein the subject is a carrier of a gene associated with increased the disorder.
  - 39. The method of claim 38, wherein the subject is a carrier of the BRCA1, BRCA2, hMSH2, hMLH1, or hMSH6 gene.

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# FIGURE 1



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Gene Description S666:9 Mas 20 protein • (42% human) RIKEN ¢DNA 4930429B21 gene	S44243 endosomal protein - human 20 %. Skivzj. skižw protein ski	insulin receptor tyrosine kinase substrate (90% human) KIP1_MOUSE DNA-PKcs interacting protein (Kinase interacting protein) (KIP) (Calcium and integrin-bin 43 %	PHZ2_MOUSE Probable oxidoreductase 0610038K03Rix 100 % get2 melanoma antigen, family L. 2	inorganic pyrophosphalase [Homo sapiens] 71 %	11a1 a1xi collagen chain	ss1 adull male hippocampus riken coma cone;2800068624 full insert sequence; deadth asp-glu-ele-aspthis box binding protein doxbp1 bbas phosphoripase c beta3		ck2 Rho-essociated colled-coll forming kinase 2  DOI 1 MOTICE Determine reliated DOI pediamental (Contains: Domese transmittens: Endominates of 12 miles due)		k? cyclin-dependent kinase ? (homotop of Xenopus MO15 cdk-activating kinase)		pp2 polyc-binding protein splice variant e: alpha-cp2 pcbp2 hrmp-e2		progesterate memorane binding protein (7.5% Monto Saplens)		ino patemany expressed efferentiation-associated-protein 2 p. 2 aanditoside-induced offerentiation-associated-protein 2		WO repeat upon and province sopherist of the	4a sodium channet, voltage-gated, type IV, alpha potypeptide	patched related protein translocated in renal cancer (28.14% Homo sapiens)		p) uoiquiin-associated protein 1		ad imprinted and ancient					נים לאומלאומותי פתח לאומלאומת פס כם לאמליכם	i3 chloride channel protein clon3	o morningo metallo androantida se mma		TLH29 profein precursor (Homo sapiens) 63 %	The state of the s	nypounencal protein FLAZOTSO (15% ilonian)		myekod cell leukemia sequence 1 2 11 days embryo riken cdna clone:2700054007; ribosomal protein rp112	
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91001		27878	75102		07073	61716	50529		73834	68942	21859		80752	35024	24127	71099	16151	18291	56709	10064	70918		69578	40242	1924	74902	58801	15980	;	71544	12986	70849	72058	27405	14626	15496	19110	2000	14455	23853	67285	321015	75356	72413	1887	7944	114875	56095	104348		93966	54139	04217	18686	į
	Cluster Access	6 BC002307	6 AK006373 6 AK015717	6 NM 008508	6 AK003611	6 NM_020520	6 AK009251	6 AK014595	6 NM 023721	6 AK004506	6 NM_011595	6 AK012931	6 NM_025482	7 NM 030565	7 AKU07 130	O COLORA C	2 MM 010547	7 AY061761	7 NM 019965	7 NM 026537	7 NM_008918	7 NM_015771	7 AK009387	7 AB047820	7 NM_016978	7 NM 023129	7 NM 021451	7 NM 008338	7 AK020725	7 AK018520	7 NM 007782	7 AK006984	7 NM 019476	7 AKU08000	7 NM 030239	7 NM 008295	7 NM_011165	7 AK011565	7 NM_02145/	7 MM 01353	7 AK014025	7 AK019938	7 AK016205	7 AK012400	7 AK020538	7 AK017110	7 AK020/39	8 NM_034000	8 NM 023266	8 AK015359	8 NM_053149	8 NM_016851	8 AK005949	8 NM_053011	8 NM USING

	torsin termity. In member 8 vaccinia refated kinase 1 vaccinia refated kinase 1 hypothetical protein FLJ23467 (Horno saplens) 93 % hypothetical protein FLJ23467 (Horno saplens) 93 % C-type (calcium dependent, carriohydrate reorganition domain) lectin, superfamily member 6 RIKEN full-length enriched Ray Jones 433405H02 zinc finger protein 34 (Horno saplens) 90% protocoather in alpha 6 protocoather in alpha 7 pro		Optioning Page, January 1, Studenting or Depression of Action (PEBP) 40 % PEBP_MOUSE Prospitation of Action (PEBP) 40 % histocompatibility 2. M region focus 2 chemokine (CC) receptor 3 evv. 2 even-skipped homeobox homatog mesodermal cell dorsoventral fates determinator this sequence comes from fg. 2 map bocation LIM domain blinding 2 Eas-activated serin-althroconine kinase (Mus musculus 31 %) paralemunin Mus musculus) 100 % unknown ephrin 81 PRP1_HUMAN Salivary profiline-rich protein precursor (Clones CP3, CP4 and CP5) (Contains: Basic peptid 28.42 % day neonate skin rich and me dome-46324 1509 RKIKEN CDNA 5/30493819 [Mus musculus] 69.05 % WO repeat domain 17 Similar to hypothetical protein FLJ11730 [97% Homos sapiens) receptor (TNFRSF)-interacting serine-threonine kinase 1 A Chain A, Crystal Structure Of Mutant Human Lysoxyme Substituted At Left-Handed Helical Positions 48 % e2t transcription factor 5 clone mgc:6043; e2f-5 protein hymbrocyte antigen 64 chromodomain protein. Y dront condendate
Gene Apobec1 Ndst4 Cd37	VACION VACIONI VALON VACION VA	Siglece Line2 Zip287 Jiph3 Jiph3 Phili Scot2 Reg2 Reg2 Nktrit Lag3	Cypton Cor3 Cor3 Cor3 Lubb2 Env2 Hsp25 Wdr17 Ripk1 E255 Ly64 Cord
Locus 11810 64580 77462 76559 12493 73937	30334 22367 69769 26888 74721 12937 1892 17391 17391 17391	54168 73732 8382 74249 170740 57340 19655 19650 17230	1307.0 14950 12771 14029 16826 69577 6836 15507 69113 244484 70088 19766 19562 77562 77562 77562 77562 77563
Cluster Access 8 NM 031159 8 NM_022565 8 AK021099 8 AK010577 8 NM 007645 8 AK015107	B AZ297743 B NIM_01705 B NIM_011899 B NIM_01899 B AK01271 B AK01271 B NIM_00994 B NIM_01898 B NIM_01899 B NIM_018997 B NIM_018998	8 NW_021569 8 AK001567 8 AK001557 8 AK010252 8 AK281141 8 AK281141 8 MG52317 8 NW_026055 8 NW_021829 9 NW_02128 9 NW_020903 9 NW_06896 9 NW_06896 9 NW_06896 9 NW_06896 9 NW_06896 9 NW_06896 9 NW_06896 9 NW_06896 9 NW_0749180 9 AK016896 9 NW_0749180 9 AK016896 9 NW_0749180	9 NM, 002954 9 NM, 002954 9 NM, 002914 9 AK01527 9 AK011281 9 NK, 01381 9 NK, 01388 9 AK01698 9 NK, 01068 9 NK, 01068 9 NK, 0108 9 AK01698 9 AK01698 9 AK01988 9 AK01988 9 AK01987 9 NM, 00799 9 NM, 00799 9 NM, 00799

	ntooline acetykonoline feetpior subunii eiphaa nicas Biren Anni Anasasasas	KIZ TO THE STATE OF THE STATE O	859254 mysoin heavy chain 12, splice form2 - human 29 %	Motoney feukemia virus 10-like 1 [Mus muscukus] 37.37	UDP-GatbetaGkCNAc beta 1.4-galactosyframsferase, potypeptide 1	Intreceptor 835ocialed factor 4	integrated to the control of the con	forther a general plant entering formally managed as	BUILD OF THE SECURITY OF THE S	homeodomain profession or something	Ferritory of the state of the s	RIKEN CYNE ASHIOLEGUE 14	RIXEN CON ACCORDING ACCORDING	denin cialmhorathan	RIKEN CHONA BASION CHONA	VAV. TELEVISION OF THE PRINCIPLE OF THE	Philosophia central ce	\$12207 hypothetical profess (RR elament) - process (RR hypothetical profess (RR hypothetical pro	Nsahasahajase 1	aduli male testis riken cdna clone: 1700048h20; clone: 4933432e 16	RIKEN CDNA D730001C10 gene	KIAA1074 protein [Homo sapiens] 45.33 %	5-hydroxythydamine (serolamin) receptor 58	ollactory receptor MOR32-4 - odorani receptor S48 gene (Mus muscutus) 97 %	FYVE and coiled-coil domain containing 1	carrello-like 2	flavin containing monooxygenase fmo3		CGI-30 protein [Homo sapiens] 85 % /	ed2 oncogene		hitherin-like arolein i		ephequin	RIKEN CONA 0610011D08 pene		RIKEN CDNA 2810030N09 gene		Mus musculus 0 day neonate skin cDNA, RIYEN full-langth enriched library, clone; 4632413E21 product; wealty similar to PHOSPHOLIPASE B (Rattus n	c-tos celular homotog to viral oncogene c-tos protein	cal eye syndrome chromosome region, candidate 6 homolog (human)	RIKEN CDNA 4930524,08 gene		Cytorrone Paso, same X, Soutiers X, Doylers 40	71 endi usama enama enama 17.	reparaments caramentarian and and an analysis of the caramentarian and an analysis of the caramentarian and and an analysis of the caramentarian analysis of the caramentarian and an analysis of the caramentarian and an analysis of the caramentarian and an analysis of the caramentarian analysis of the caramentarian and an analysis of the caram	SUSPENDENT FINEN CLINA SUCCESSED GENE	DIRECT PRINT POLICE TO	RINEM SURVA SO INGOSOU BOTTO	the design of the following the first the following the first the following the first	oppose parameters of the second parameters of	ADJUGOVITE TOUR STATES		myelodysplasia/myeloid leukemia factor mif1; hematopoietic lineaga switch 7 his7	4300 protein aa 1-77		brancheding media El 1990 History envisors DO M	npocence proces in Landon I and Marie Supers (or Milenan enriched librer), done 4921520P21 product incohetical Microbodies C-terminal terreting terminal terreting terret	
E 2	Curna	Kir21b			B4gall I	1014	Roll 30	Cohen	5	5	Xrt1-14			Demo		Vava	Wise		Lyplat	:			HIS	MOR32-4	Fycol	CHI	Fmo3			Ect2		Tudio t	•	Ereg	)					8	9 2 2 3		0	Cyprosi	7166		700.003	70107		144	2	Ads		MIT	Hove2				
Locus	75737	16565	74717	71159	14595	13040	59044	57249	71469	12951	16654	212127	75287	13517	77849	22325	67712	71546	18777	67003	78589	70952	15564	258755	17281	93673	14262		69740	13605		56784		13874	66588		69259		78261	14281	94047	75129	13000	2000	2000	00000	1000	72015	70240	22045	102204	11875		17349	15456		87504	70887	
Cluster Access		10 NM 019962	10 AK015780	10 AK016833	10 NM 022305	10 NM 003423	10 NM 021540	10 NM O2048B	10 AKD18420	10 NM 007770	10 M13806	10 AK012908	10 AK016085	00010 MN 01	10 AK021026	009500 MN 01	10 AK019700	10 AK018561	10 NM 008866	10 AKO06727		10 AK015473	10 NM_010483	10 NM_020292	10 AK0 13060	10 NM_053096	10 NM_008030	11 NM_019540	11 AK010475	11 NM 007900	11 NM 026204	11 AK009111	11 AF121976	11 NM 007950	11 NM 025647	11 D29987	11 AK011637	11 AK005181	11 AK019494	11 NM 010234	11 NM 033567	11 AK015886	11 AK005050	11 AM 080454	T NIM USUASA	11 AKO10406	00 MM 01250	44 AKO14453	14 AF312033	11 NM 013696	11 ARD-54457	11 Y08028	11 AB041591	11 NM_010801	11 NM_008283	11 NM_008085	9		

Description	RIKEN cDMA 1700006C06 [Mus musculus] 100 %	G-protein-coupled receptor 50	cholinergic receptor muscarinic 4 chrm4	Usher syndrome 1C homolog (human)	meprin a alpha-subunit mep-1	RIKEN cDNA 1110034G24 gene	Mus musculus mRNA for CN 8 soFv. complete cds	dolichyl-phosphate alpha-n-acetylghcosaminephosphotransferase 2 dnautz	cyclin a 1	RIKEN cDNA 2310041H08 pene	polymerase (DNA directed), arbha 1	RIKEN CONA 2310016N21 IMus muscubal 100 %	polassium large conductance calcium-activated channel subfamily M ainha member a	chondroitin 4-sulfolransferase [Mus muscutus] 47 %	- 138467 testin - human 51.90 %		uncharacterized hematopoletic stembrogenitor cets protein MDS031 (Homo sepiens) 86 %	teroid 5 atotra-reductase 2-tike; HSAR gene; steroid 5 atotra-reductase 2 like [Mus muscutus] 100 %	nuclear pore membrane dycopratein 210 pom210	13 days embryo stomach riken cdna clone:d530014Q21	zinc finger protein 142	secreted frizzled-related sequence protein 5	fibroblast growth factor 3	melanocyte prolifeating gene 1	TATA box binding protein (Tbp)-associated factor, RNA polymerase I. A	casein delta	solute carrier famity 39 (zinc transporter), member 4 (Homo sapiens) 39 %	pir:S34338 - S34338 billary glycoprotein F - mouse 34 %
Gere		Gpr50	Chime	Ushtc	Mep 1a			Dpag12	Son		Polat		Konmal					Srd5a2l	Nup210		Zfp142	Strp5	For	Myg	Tafle	Csnd		
Locus	66320	14765	12872	72088	17287	73747		13478	12427	75686	18968	75581	16531	71797	78733		67574	57357	54563	78717	77264	54612	14174	60315	21339	12992	72002	71601
Cluster Access	11 NM_025486 11 NM_053188	11 NM_010340			11 NM_008585	12 AK004090	12 AB036341	12 NM_007875	12 NM_007628	12 NM_029385	12 NM_008892	12 NM_023784	12 NM_010610	12 AK004401	12. AK021408	12 AB041649	12 NM_026247	12 NM_020611	12 NM_018815	12 AK021287	12 AK020384	12 NM_018780	12 NM 008007	12 AK017955	12 NM_021468	12 NM_009973	12 AK007473	12 AK018613

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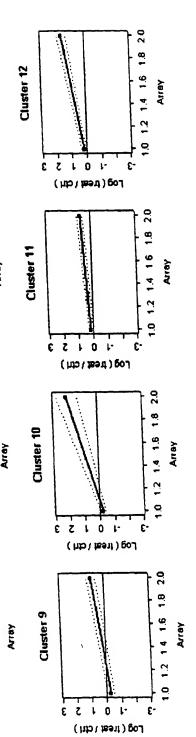
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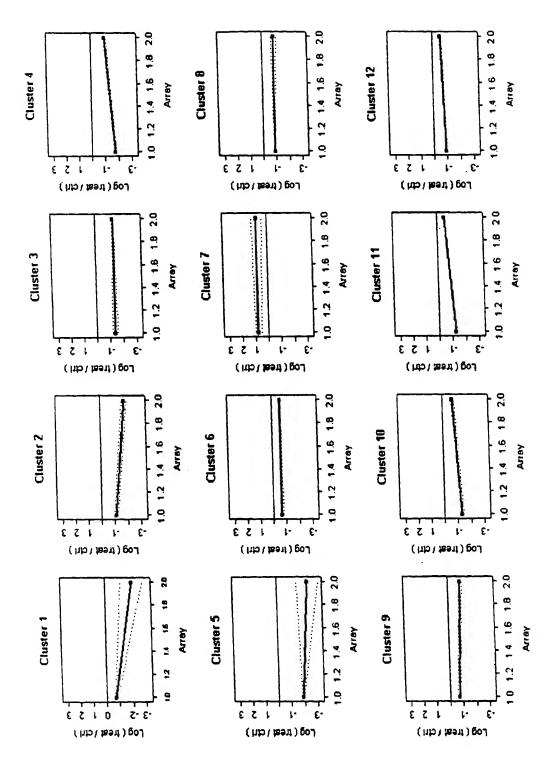
Cluster 4

Cluster 8

Array



Description	DNA segment, Chr 8, ERATO Doi 633, expressed	offactory receptor 154	matrilin 4	syntaxin binding protein 4	gooseoid gsc	Mus musculus RIKEN cDNA 2410116G06 gene (2410116G06Rik)	C-teminal PDZ domain ligand of neuronal nitric oxide synthase	calsequestrin 2	GPI-anchored metastasis-associated protein homotoo	anistatess 4	protein phosphatase, EF hand calcium-binding domain 2	glia maturation factor, gamma	Co2+-dependent endoplasmic reticutum nucleoside diphosphatase	RIKEN dDNA 4933406P04 pene	fibroblast activation protein	S12207 hypothelical protein (B2 element) - (80% mouse)	ref:NP 061935.1 - hypothetical protein FLJ20225 [Homo sapiens] 68 %	Mus musculus 6 days neonate head cDNA, RIKEN full-length emidted library, clone:5430417C01 product hypothetical profein	growth differentiation factor-9b gdf-9b; bone morphogenetic protein 15 bmp15	ructear factor of kappa light potypeptide gene enhancer in B-cells inhibitor-like 1	phosphodiesterase 7A	protein tyrosine phosphatase, non-eceptor type 12	hypotheticat protein FLJ20473 (Homo sepiens) 69 %	DNA segment, Chr. 1, ERATO Doi 396, expressed - RIKEN cDNA 2610307121; hypothetical protein, MNCb-4273 [Mus musculus] 100 %	tumor-suppressing subchromosomal transferable fragment 3	Similar to sperm adhesion molecule 1 (PH-20 hyaturonidase, zone pelitucida binding) [45% Homo sapiens]	Similar to CUB and Sushi multiple domains 1 I74% Mus musculus	sodium channel, voltage-gated, type VI, alpha polypeptide	Similar to A40038 MHC class I histocompatibility antigen H-2 M3 alpha chain precursor - mouse 62%	RIKEN CDNA 4931431F19 gene		RIKEN cDNA 2810206C17 gene	phosphodiesterase 1C	rhodopsin kinase	JC6547 high sulfur protein B2E - rat 37 %	neurogranin	interferon regulatory factor-3 irl3 irl-3; factor 3	RIKEN cDNA 1700006J14 gene	cDNA 1810008A14 [Mus musculus]	major vault protein	niemann pick type c1
Locus Gene	52447	27216 Ollr154	17183 Matn4	20913 Stybp4	14836 Gsc	68236	70729 Capon	12373 Casq2	72434 C4 4a	11695 Alx4	19023 Ppef2	63986 Gmfg	76025 Shapy	74420	14089 Fap	77252	66825	78548	12155 Bmp15	18038 NRbil1	18583 Pde7a	19248 Ptpn12	71699	52477	22113 Tssc3	77042	74943	20272 Scn7a	76797	70980	69171	72495	18575 Pde1c	24013 Rhok	96969	64011 nrgn	54131 1/13	321010	66268	78388 Mvp	18145 Npc1
Cluster Access	1 AROLDS	1 NM 013/28	1 NM_013592	1 NM_011505	2 NM_010351	Z NM_026630	2 AK018149	2 NM_009814	3 BC016549	3 NM_007442	3 NM_011148	3 NM_022024	4 AK006565	4 AK016707	4 NM_007986	4 AK020460	5 AK020617	5 AK019946	6 NM_009757	6 NM_010909	6 NM_008802	6 NM_011203	6 BC011108	7 NM_021421	7 NM_009434	8 AK014599	8 AK015672	8 L36179	8 AK010800	8 AK016497	8 AK007667	9 AK011897	9 NM_011054	10 NM_011881	10 NM_027170	11 NM_022029	11 NM_016849	11 AK005678	12 NM_025457	12 NM_080638	12 NM_008720



Description	catabolite repressor protein ccr4-associative factor caf1; cc4-not transcription complex subunit 7 done mgc:6050	ref:NP_478062.1 - chromosome 21 open reading frame 51 [Homo sapiens] 82 % hypothetical protein FLJ20758 (73% human)	adult male testis riken cdna clone:4930348g07, 4930370g07, adult male testis riken cdna clone:4930348g07, 4930370g07, 493070g07, 493070000g07, 493070g07, 493070g07, 493070g07, 493070g07, 493070000000	v-ral sinian learning train oncogene hornolog B (ras related) v-ral sinian learning oncogene hornolog B (ras related)	ENCT MOUSE Economics of the property of the pr	RIKEN CDNA 1700084J12 gene	discs, targe homolog 1 (Drosophila) ORM1-like 3 (S. cerevisiae)	in the second state of the second sec	steroidogemic acute regulatory processions and statement of the statement	AFG3(ATPase family gene 3)-like 1 (yeast		ubiquilin-specific protease otubain 2 (94%)		catechol-o-methyltransferase comt			75-q1z	opioid grown factor recopied	ffc, ffrigetine i (co-co-co-co-co-co-co-co-co-co-co-co-co-c	HSPC134 protein [60% Homo sapiens]	karyopherin (importin) alpha 6	sterol regulatory element binding protein srebp1; clone image:3590844			homeo box 88	
Gene	Cnot7		;	Mecp2 Raib	646	CIEK	Digh1		MIn64	Afg3l1				Com	Mors1		Zfp57	Ogt	Frd1	B3gallZ	Kona6		10310		Hoyb8	-
Locus	18983	68936	69950 75339	17257 64143	76553	50883 73486	13383	71999	59045			9	6+100	12846	51812		22715	72075	76157	26878	75608		2078/		9	0.00
Access 1 L29479	1 NM_053188 1 NM_018779 1 NM_019709 • NM_011135	1 NM_024217 1 AY033901	1 AK013131 1 NM 023773	1 NM_010788	1 AK012967	1 NM_016681 1 AKOO6993	1 AK016641 1 NM_007862	1 NM_025661	1 NM_021547	1 AK008844 1 AK008844	1 NM_020563	1 AK014242 1 BC004639	1 NM_026580 1 AF217002	1 NM_021471	1 NM 016766	1 AK006388 1 NM_013562	1 AK005168	1 NM 00353	1 AK018094	1 NM 020025	1 BC006905	1 NM_008456 1 AK016603	1 BC006051	1 NM_021405	1 NM_025980	1 X13721 1 NM 023805
Cluster																										

e Description e prolactin-like protein E 16 zinc finger protein 146 zfp146 2 inc finger protein 146 zfp146 2 incurrophil cytosolic factor 2 3 otogelin mucin-like extracellular matrix protein mlemp 3 ribosomal protein s6 rps6	Mus musculus mitochondrlon, complete genome  a actin related protein 2/3 complex, subunit 3  RIKEN cDNA 2410019A14 gene thyrotrophin-releasing hormone receptor th-r; thyrotropin releasing thr  3 small nuclear ribonucleoprotein D3		Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:4921520P21 product:hypothetical Microbodies C. solute carrier family 6 neurotransmitter transporter faurine member sloca6; mus cookii taurine/beta-alanine leptin PTK9 notein krosine kinase 9.	CpG binding protein  AMRP_MOUSE Alpha-2-macroglobulin receptor-associated protein precursor (Alpha-2-MRAP) (Low density I 99.72 % adult male testis riken cdna clone:1700122011  CD24a antigen
Gene Pripe Zfp146 Ncf2 Otog Rps6	mt-Nd6 Arpc3 Trhr Snrpd3	Ssbp3 Mag Plscr2	Sic6a6 Adar3 Lep	Cgbp Lrpap1 Cd24a
Locus 19113 26465 17970 18419 20104	17722 56378 69746 22045 67332	71020 72475 17136 18828 68175	70887 21366 94191 16846 19230	74322 16976 76651 12484
Cluster Access 1 NM_00830 1 NM_011980 1 NM_010877 1 AK005138 1 AK016624 1 NM_013624 1 NM_013624 1 NM_019794	1 NM_030724 1 NC_001569 1 NM_019824 1 AK010555 1 NM_013696 1 U56773 1 NM_008040	2 NM_021394 2 AK005865 2 AK011853 2 AK010425 2 NM_010758 2 NM_008880 2 NM_026596 2 NM_026596	2 AK014894 2 NM 019945 2 NM 025826 2 L10319 2 AK012521 2 NM 025774 2 NM 005320 2 NM 006320 2 NM 006330 2 NM 006330	2 NM_009745 2 NM_01960 2 AK010337 2 S67967 3 AK018071 3 AK007241 3 NM_023480 3 AB039178 3 NM_009846 3 NM_009846

Description hypothetical protein FLJ10199 [84% Homo sapiens] hypothetical protein FLJ10199 [84% Homo sapiens] RIKEN cDNA 1700034J05 gene cellular retinoic acid binding protein 1 nedd4 ww domain-binding protein 5 n4wbp5 leukocyte surface antigen cd37	heat shock factor hsf2 ras-related protein rab18	bystin-like	solute carrier family 25 mitochondrial carrier ornithine transporter inclinacy. The property of the property o	prostaglandin D receptor enguity 2, ced-12 homolog (C. elegans)	RIKEN cDNA 2410002022 gene RIKEN cDNA 800491400040141 membrane 23 homolog (yeast)	translocase of inner mitoducing as the property of the propert	point in process on cogene family 9	Mus musculus adult male stomach CDNA, Kirken ruin-tengui emismo.	RIKEN CDNA 5/304855 9 (22.5) 7	prain neurotensiin coopers a regional praintensiin coopers a r	RIKEN CDNA 4930542N07 gene	paraxis basic-fletix-toop norm programmer and the paraxis basic-fletis riken of the done:1190010!23 adult male testis riken of the done:1190010!23	RIKEN cDNA 493140/Nuz gene hras suppressor expressed in skeletal muscle heart brain and bone marrow; 18 days emusyoning 40.00 %	teucine-rich and death domain containing: p53 protein induced, with deam domain times moscored.	RIKEN CDNA 4933407G07 gene Ketohexokinase (Hepatic fructokinase)	vesicle-associated membrane protein ceinure in 1977. SYN1_MOUSE Synapsin I 30 %	histone deacetylase hdac7 adp-ribosylation factor 5 arf5	gcn5 histone acetyltransferase; clone Image:3491089
Gene Crabp1 N4wbp5 Cd37	Hsf2 Rab18	Bysl	Sic25a15	Ptgdr Elmo2		Timm23 Gsh1		Kappa	Zfp371	Ntsr2 Gzma		Tc/15	Hrasrs		ž	Vamp3	Hdac7a Arf5	Gcn512
Locus 214359 73344 12903 65113 12493	15500 19330	53414	18408	19214	66975	53600	13388	19340 Unknown 233802	74476	18217 14938	75185	21407	77627	70388	71062	22319 69539	56233	14534
<b>Čiuster Access</b> 3 BC003277 3 AK006598 3 NM_013496 3 AF220209 3 AF20039	3 NM 031402 3 NM 008297 3 NM 011225	3 AK005878 3 NM_016859	3 AK005902 3 NM_011017	3 AK006440 3 NM_008962	3 NM_060267 3 AK012141	3 AND 016897 3 NM 016897 3 NM 008178	3 NM_007865 3 AK010654	3 NM 031874 3 AK019053	3 BC004775 3 AK017115 3 AE904778		3 NM_013818	3 NM_009328	3 AKU07246 3 AK016516 3 AM 013751	3 MM_015751 3 AK016461 3 AK008713	3 A8023957 3 AK016718	3 NM_008439 3 NM_009498 3 AKOOOO4	3 NM 031881 3 NM 019572 3 NM 007480	3 NM 013738 3 NM 020004

Description	expressed sequence AW123240	CRYL_HUMAN Lambda-crystallin homolog 83 %	T-cell leukemia, homeobox 2 RaiBP1 associated Eps domain containing protein	PKE protein kinase, hypothetical serine/threonine protein kinase [Mus musculus] 100 % KLKC_HUMAN Kallikrein 12 precursor (Kallikrein-like protein 5) (KLK-L5) 70 %	inducible T-cell co-stimulator S43344 sex-determining protein Sry - mouse (27 % M. musculus)	myspdz myosin containing pdz domain hypothetical protein FLJ20772 (79% human) putative cerebral cortex transcriptional regulator t-brain-1 tbr-1 regulator t-box brain tes-56 tocopherol (alpha) transfer protein trioartite motif protein 17	enolase 2 gamma neuronal eno2	RIKEN CDNA 1700026D08 gene	rhotekin expressed sequence Al604832 keratin complex 2, gene 6g [Mus musculus] 100 % multiple PDZ domain protein	Tax interaction protein 1 [Homo sapiens] 99 % hepatocellular cardinoma-associated antigen 112 [55% Homo sapiens] hepatocellular cardinoma-associated antigen 112 [55% Homo sapiens] phospholipase A2, group X immunoglobulin superfamily containing leucine-rich repeat immunoglobulin superfamily containing leucine-rich repeat RIKEN cDNA 5830411J07 gene
Gene		Cryl1	Hox1111 Reps1	Pke	Icos	Tiaf1 Tbr1 Tips	Eno2		Rtkn Krt2-6g Modz	Pla2g10 Isfr
Locus	101333	68631	21909	57740	54167	21842 68260 21375 50500	13807	75556	20166 102566 56735	76281 66058 26565 26968 74742
	3 S82853 3 AF071068 3 D64112 3 AK009770 3 NM_053201		3 AK020683 3 AK019434 3 BC005625 3 NM_009392 3 NM_053258	3 NM_009046 3 BC005556 3 AK006258 3 NM_021302 9 AK009213	3 AK019591 3 NM 017480	3 AKUZU613 3 AF343088 3 AB02647 4 AK019481 4 NM_009322 4 AK004882	4 AF220135 4 AK007760 4 AK021211 4 ARC03842	4 NM_030722 4 AK006375	4 NM_025746 4 NM_009106 4 BC002294 4 NM_019956	4 NM_010820 4 Z67747 4 AK004963 4 AK002546 4 AF210429 4 NM_012043 4 NM_029082 4 AK016622 4 BC002120

Description	il-3 receptor alpha subunit	actinin alpha 3	RIKEN cDNA 1700013G24 gene Mus musculus 13 days embryo male testis cDNA, RIKEN full-length enriched library, clone:6030449J2U product.rypostation	histone hat hist putative; testicular ha	purine-selective na+ nucleoside cotransporter cntz cntz spin cn	protyl 4-hydroxylase alphai-subunit	wee 1 homolog (3. During) Mus musculus adult male testis cDNA, RIKEN full-length enriched library, conc. 1000		RIKEN CDNA 9230112K08 gene	RIKEN CONA 191905. Similar to A56465 transcription factor IIIC2 beta chain - human oo 76 Similar to A56465 transcription factor IIIC2 beta chain - human oo 76	tachykinin receptor 2 tacr2	S50853 translation releasing factor every (1997) in the control of	Ellis van Creveld gene nomotog (numen.)	A55253 Instance	cytocal oring Control of the Control	RUZA_HUMAN UZ small nudear ribonuceoprocess	methionine aminopephoase-line (			Similar to ROSIT [54% Rattus norvegious]		hypothetical protein FLJ21415 (93% numeri)	mrpig mitochondriai ribosorinai processi	arec3	transcription elongation regulator (CTCO) (85.88% Homo sapiens)	hypothetical protein FLJ20156 (Homo sapletis) 20 %	RIKEN CUNA O' 2010 11 80 11		
Gene	II3ra	Actn3		H1f3	Slc28a2	P4ha1	Wee1				Town	1907	Evc		Cyp11b2		Metapl1	Abpl	Cdc42ep4				Mrp <sup>19</sup>	SivA	Tcerg1				
Locus	16188	11474	69380	68148 14957	59023	18451	22390	1761	78081	69186		21337	59056	77836	13072	14466	66559	76507	56699	74057		76702	78523		56070	70544	77891		
Chister Access	4 AK016374 4 NM 022813 4 NM 008369	4 AK016436 4 NM 013456		4 AK020067	4 AK015845 4 NM 021520	4 NM_030703 4 AK013485	4 U16162 4 NM_009516	5 AK016155 5 NM 023737	5 NM_026631	5 AK007623 5 AK007623	5 NM 018873	5 NM_009314	5 BC013717	5 NM 02 25 25 25 25 25 25 25 25 25 25 25 25 25		5 AF127033	5 AK016859	5 NM_U25633	5 NM 020006	5 AK016661	5 AK014544 5 AK005345	5 AK015708	5 AK010791 5 AB049637	5 NM 021328	5 D50416 5 NM 019512	5 AK017626	5 AK019545 5 BC012255	5 NM 021713	S NM OCOCA

## FIGURE 4-6

Description ethanol incuded 2 [putative ribonuclease III; putative protein p241 which interacts with transcription factor Sp1 [Hom 99 % ] hypothetical protein FLJ12806 [Homo sapiens] 97 % / 305 aa interferon alpha family, gene 4 adult male testis riken cdna clone:1700001002 RIKEN cDNA 4930544L04 gene	A Chain A, Glutathione Transferase (62% human) solute carrier family 22 (organic cation transporter), member 9	Mus musculus odorant receptor M15 gene keratin complex 1, acidic, gene 17 S100 protein, beta polypeptide, neural catalase per-hexamer repeat gene 1	high mobility group protein i isoform c hmglc zinc finger protein Similar to vesicle transport-related protein [95% Homo sapiens]	intronless potassium channel lint i chemokine (C-C motif) ligand 2	galnac alpha-26-sialyltransferase i	winged-helix protein twh transcription factor; forkhead box b1b foxb1b CGI-74 protein; CGI-59 protein [Homo sapiens] 98 % cholinergic receptor, nicotinic, beta polypeptide 2 (neuronal) RIKEN CDNA 4833426110 gene	F-box and WD-40 domain protein 7, archipelago homolog miwi piwi piwi biwi piwi DNA (cytosine-5)-methyltransferase 3A (Dnmt3a) (DNA methyltransferase MmulliA) (DNA MTase 30 % DM3A_MOUSE DNA (cytosine-5)-methyltransferase 3A (Dnmt3a) (DNA methyltransferase MmulliA) (DNA MTase 30 % 2-hydroxyphytanoyl-CoA lyase tyrosinase related protein lyrp1	target of myb1 homolog (chicken) dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 isoform 2 [99% Homo saplens] gamma-aminobutyric acid (GABA-A) transporter gamma-aminobutyric acid (GABA-A) transporter F-box only protein 8b
Gene Etchi2 Ifna4	SIC22a9	Krt1-17 S100b Cas1 Phxr1	Hmga2 Zfp161	Kcna1 Ccl2	Siat7a	Foxb1 Chmb2	Fbxw7 Piwil1 Hpcl Tyrp1	Tom1 Gabt1 Fbxo6b
Locus 14000 236519 15967 75435	68214	20203 12359	15364 22666 76983	16485	20445	64290 192196 11444 74592	50754 57749 70802 56794 22178	21968 74415 232333 50762
Cluster Access 5 AK003651 5 BC004835 5 NM_010504 5 AK005616 5 NM_007411 5 AK019737	5 AK014696 6 NM 023502 6 AK018953 6 NM 023154	6 NM_019723 6 NM_025325 6 AF282300 6 NM_010663 6 NM_009115 6 NM_0099145	6 NM 011000 6 AK016590 6 NM 010441 6 NM 009547 6 AK010070		6 D29987 6 AK016572 6 NM 011371 6 AK007262		6 AKU14112 6 NM_026187 6 NM_080428 6 NM_021311 6 AKU14534 6 NM_0319975	6 NM_026125 6 NM_011622 6 AK016565 6 L32178 6 NM_015797

e Description 55 mitochondrial ribosomal protein S5	3 connexin31 f2 POU domain, class 3, transcription factor 2 a atonal homolog 7 (Drosophila)		cyclin-dependent kinase-like 2 (CDC2-related kinase)	cyclin h ccnh regulatory subunit of cdk7 partner cdk7 in cak and tfiih BTB (POZ) domain containing 1 [Homo sapiens] 26.88 %	insulin receptor tyrosine klnase substrate (90% human)	expressed in hematopoietic cells, heart, liver adult male kidney riken odna clone:0610027d24	12 days embryo embryonic body between diaphragm region and neck riken cona clone:9430077c05 11 SSN1_MOUSE SAM-domain protein SAMSN-1 (SAM domain, SH3 domain and nuclear localisation signals prote 100 % RIKEN cDNA 4833420G17 gene	ecyt-Coenzyme A oxidase 2, branched chain splicing factor 3a, subunit 1 RIKEN cDNA 1200014H24 [Mus musculus] 100 %		lymphocyte antigen 86 cut drosophila-like 2 cutl2	<ul> <li>translocase of inner mitochondrial membrane 8 homolog a yeast clone mgc:6730; timm8a</li> </ul>	squalene synthase ii ia antigen-associated invariant chains ii31 and ii41 exons - 8 protein; done mgc:6517	BCL.2/adenovirus E1B 19kDa-interacting protein 3-like zinc finger protein 295 (32% human)	RIKEN cDNA 1500031J01 gene sry-box containing 7 sox7 midline 1	histone deacetylase 5 hdac5 dass ii
Gene Mrps5	Gjb3 Pou3f2 Atoh7	Kcnj2	Cdki2	Ccnh		Hhi Biod2	Samsn1	Acox2 Sf3a1	Osbpl3	Ly86 Cutt2	Timm8a	Fdft1 Ii	Bnip31	Sox7 Mid1	Hdac5
Locus 77721	14620 18992 53404 17913	16518 69347	53886	66671 224671	66898	29809 76895	77297 67742 67392	93732 67465	71720	17084 13048	30058	14137 16149	12177 75580	67529 20680 17318	15184
Cluster Access 6 NM_029963	6 NM_008126 6 NM_008899 6 NM_018664 6 NM_018664		6 NM_025529 6 NM_016912 6 NM_026645 6 RC003372	6 NM_023243 6 AK006275 6 AK006092			6 AK020491 6 NM_023380 6 AK014735	6 NM_053115 6 NM_026175 6 NM_023803	6 AK004768 6 BC004690	6 NM_010745 6 NM_007804 6 NM_007485	6 NM_013898 6 AK01447 7 NM 019541	7 NM_010191 7 BC003476 7 AK007368	7 NM_009761 7 AK009517 7 1 26164	7 NM_026218 7 NM_011446 7 NM_010797 7 NM_025597	7 NM_010412

	surx protein putative atpase and nelicase sarcotipin 1 dentin matrix protein 1 interfeukin 4			L1 repeat, Tf subfamily, member 29 [69% Mus musculus]	hypothetical protein FLJ10560 [82.88Homo sapiens]	c protein phosphatase 1, catalytic subunit, gamma isoform nuclear localization signals binding protein 1 (Mus musculus 79 % )	arginase type II	adult male kidney riken cdna clone:0610011i20; deiodinase iodothyronine type i dio1 RIKEN cDNA 2410157M17 gene		prolactin regulatory element binding	10 days neonate skin riken odna done:4733401i23 k ubiquitin specific protease 27, X chromosome	oxysteral binding protein-like 5	hyaluronan synthase 3 has3 Similar to T14765 hypothetical protein DKFZp434N014.1 - human (fragment) 95 %		INDESIT	v-maf musculoaponeurotic fibrosarcoma oncogene family protein k avian mafk; erythrold transcription factor nf-e2 subunit	-		SET translocation [Mus musculus] 100 %	mudix (nucleotide diphosphate linked motely X)-type motif 3	ADP-ribosylauon-like 4
Gene Pola1	Sin Omp1		Baiap1			Ppp1cc	Arg2	Dio1	Semcap2	Preb	Usp27x	Osbp15	Has3	,	MSM	Mafk	Cyp2u1	Hsd3b5	Set	Nudt3	Ark 4
Locus 18968 55950	22389 66402 13406 16189	269846	14924	75110	70573	19047 74125	11847	13370 76820	54120	20907	Unknown 54651	79196	15118 73750		969/1	17135	71519	15496	56086	56409	11861
Access 7 NM_015819 7 NM_008892 7 NM_018772	7 NM_025540 7 NM_025540 7 AJ242625 7 NM_021283 7 AK004845	7 AE000664 7 NM_019629 7 NM_025389	7 AF027505 7 NM_023697	7 AK015778 7 NM_010016 7 NM_053156	7 AK017705 7 NM_021493	7 NM_013636 7 AK004793 7 BC003852	7 NM_009705 7 AK017416	7 NM_007860 7 AK010821	7 NM_016867 7 BC004761 7 AK021330	7 NM 016703 7 AK013879	7 AK014638 7 AF229643	7 NM_024289 7 AK007061	NM_008217 AK004110	NM_007935	NM_011304		AK018458	NM 008295	NM_023871	NM_019837	NM_007487 NM_007725
Cluster			•		14	10 10	-		~ <b>~ ~</b>	~ ~	~ ~	~ ~	~ ~	~ 1	. ~	~ 1	~ ~	^	7	<b>^</b> 1	· /-

Description	RIKEN cDNA 2410030K01 gene recombining binding protein suppressor of hairless-like (Drosophila)	CGI-99 protein [Homo sapiens] 97 % RIKEN cDNA 1700055O19	phospholipase C, zeta 1 PHD zinc finger protein XAP135, isoform a [Homo saplens] 96 %	transgene insert site 737, insertional mutation, polycystic kidney disease (NTPDase1) (Ecto-ATP diphosphohydrolase) 44%	END1 HUMAN Ectonucleoside triphosphate diphosphoriyu otasa 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Similar to Env. 1.2. The subunit 2 s	cleavage stimulation laws. Of protein), gamma 4 subdim	gluaring	hypothetical process:	carboxypeptidase B2 (plasma)	F-box and leucine-floring protein 4 (HSF 4) (Heat shown and leucine-floring protein 4 (HSF 4)	Jak-binding protein 1	putative glycolipid transfer process.	chemokine (C.C motif) receptor 1				microtubile-actin Grossman's 2	nuclear factor, complement component 3a coart	interferon gamma receptor 2 - a - n motetin-coupled receptor binds and response	anaphylatoxin c3a receptor incare is a manaphylatoxin c3a receptor incare is a manaphy	ISL1 transcription factor gamma	hepatocyte modern					and the polocity of the second			
Gene	Rbpsuhl		Picz1			Hixb9	Cstf2	Grig4 Cvo2b19		Afp Cobs	Fbx13a			,	3	Map2k1		Adp7	Nfe2l3	2	Imgrz Caara	Isl.	Hnf4g	Pols	Fino	Orce			Pdcd1	Rassio	2
9100	Locus 67629 19668	68045	114875	72057	21821	72090	15285 108062	14706	216549	11576	56373	71066	27374 79554		12768	26395		44.426	18025	73166	15980	1226/	30942		14261	56452	73707		18566	73246	56807
		8 NM_009030 8 NM_019701 8 NM_026528		8 NM 024250 9 NM 023219	8 NM_009376	8 AK003981 8 AK008590	8 NM_019944	8 AF317332 8 NM 010317	8 NM_007814	8 BC004650 8 NM 007423	8 NM 019775	8 AF176521 8 AK016803	8 AF167573	8 NM_024472 9 AK005253	8 NM 009912	8 AK009137	8 AK009704	8 AF435852	8 AF150755	8 NM_010303	8 BC01238	8 NM_009779	8 NM 021459	8 NM_013920 8 AK005864	8 NM 025396	8 NM_010231	8 NM 019710 8 AK010727	8 AK014545	8 BC006045	8 NM_006735	8 NM 020270

## FIGURE 4-11

Gane Description			74665 RIKEN cDNA 4930449E07 gene	16204 Fabo fativacid binding prolein 6. ileal (gastrotroph)			Cry1	Uble1a	55981 Pigb phosphatidylinositol glycan class b	,	l cerg1	2/413 Accol Hit-bring Sassette, Suc-tamping (MDK/IAP), member 11	occurrence of control of the control		11535 Adm preproadrenomedullin; adrenomedullin adm	Smst	Slc30a6	17423 Ndst2 glucosaminyt n-deacetylase / n-sulfotransferase dual enzyme activities; heparan sulfate n-deacetylase/n-sulfotransferase		;	15551 Htr1b 5-hydroxydryptamine (serotonin) receptor 18			52064 DNA segment, Chr 5, ERATO Doi 33, expressed	70114 FRIH_MOUSE Ferritin heavy chain (Ferritin H subunit) 41 %	70885 hypothetical protein FLJ10569 [100% human]		73731 RIKEN cDNA 1110001M24 gene	19733 Rgn regucaldin	,			14829 Grpr gastrin releasing peptide receptor grpr	12192 Zfp36i1 zinc finger protein 36, C3H type-like 1		Rps7	13849 Ephx1 epoxide hydrolase 1, microsomal	
Cluster Access	0	9 AK017352	9 AK015430 9 AK007964		9 AK002512	9 NM_018819	9 NM_030004	9 NM_019748	9 NM 018889	9 ANO17720	21280 MM 8	9 AVOA121			9 NM 009627		9 BC005753	9 NM_010811	9 NM_023908				9 AK014093		9 AK008151	9 AK014939	9 AK015898	9 AK003253	090600 WN 6	10 AK014128	10 NM 024291	10 NM_025462	10 NM_008177	10 NM_007564	10 AK016695	10 NM_011300	10 NM_010145	10 AK019535

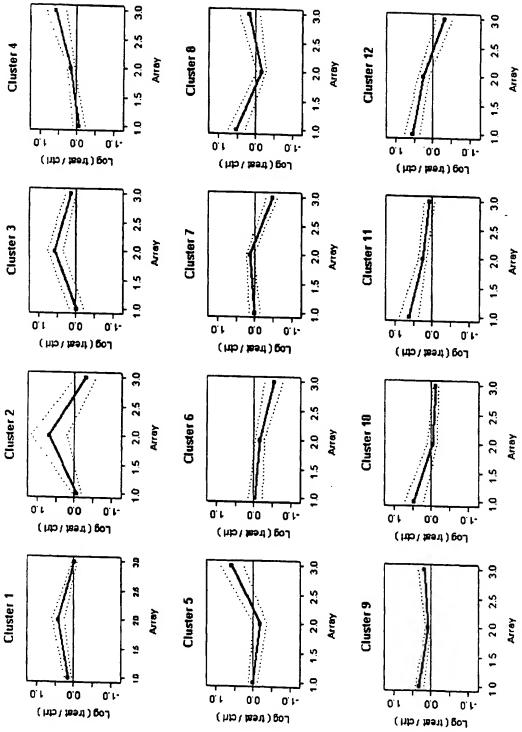
Description CGI-130 protein [71% Homo sapiens] dentatorubral pallidotuysian atrophy p21 (CDKN1A)-activated kinase 4 RIKEN cDNA 2010003119 gene	CGI-127: yippee protein [100% Human] oocyte specific homeobox 1 dine metallopeptidase damage-induced neuronal endopeptidase Similar to ISHUSS protein disuffde-Isomerase (EC 5.3.4.1) precursor - human 31 % Ariadne protein RIKEN cDNA 1810011K17 gene small inducible cytokine a28 scya28	ubiquitin specific protease 20 DnaJ (Hsp40) homolog, subfamily C, member 3 sterol-c5-desaturase potassium inwardly-rectifying channel, subfamily J, member 8 chemokine (C-C motif) ligand 19 Mus musculus gene for Cox17p, complete cds mitogen-activated protein kinase kinase kinase 4 RIKEN cDNA 2810405F18 gene Similar to apoptotic chromatin condensation inducer in the nucleus [93% Mus musculus] U2 small nuclear ribonucleoprotein auxiliary factor (U2AF), related sequence 2 T00260 hypothetical protein KIAA0605 - human 97 % sestin 2	limb-bud and heart F11 receptor RIKEN cDNA 49304290.20 gene non-catalytic region of tyrosine kinase adaptor protein 2 growth factor receptor bound protein 14 25-hydroxyvitamin d3 1alpha-hydroxylase gene for odorant receptor A16	synapsin ib syni synaplic veside associated protein RIKEN cDNA 1110014N07 gene, nuclear RNA polymerase I small specific subunit [Mus musculus] 100 % collagen a1v
Gene Drpla Pak4	Obox1 Ecel1 Arih1 Scya28	Usp20 Dnajc3 Sc5d Kcrij8 Cd19 Map4k4 Uzaf1-rs2	Lbh F11r NGK2 Gr014 Cyp40 Ora16	Syn1 Col5a1
Locus 69692 13498 70584 69870	69687 71468 13599 71830 23806 226591 56838	74270 19107 235293 16523 24047 26921 772181 75913 22184 77794	77889 16456 74626 17974 50915 13115	20964 66136 12831
Cluster Access 10 AK009957 10 NM_007881 10 AK017713 10 AK008069 10 AK007167	10 AK010201 10 AK018362 10 NM_021306 10 AK005692 10 AJ130977 10 AK008013 10 BC002098 10 NM_020279	10 NM_031387 10 AK006800 10 NM_008929 10 AB016248 10 NM_008428 10 NM_011888 10 AR047323 10 AK012994 10 AK016319 10 NM_009453 10 AK020831	10 AF317517 10 U89915 10 AK015245 10 NM_016249 10 NM_016719 10 AB006034 10 AK010153 10 AK002371 10 NM_020515 10 NM_010783	10 NM_013680 10 AK014178 10 NM_023162 10 AK016792 10 NM_015734 10 NM_026167 10 AK018093 10 NM_007863

FIGURE 4-13

Cluster	Access	Locus	Gene	Description
•	10 BC003217			
_	10 NM_016709			
-	10 NM 025476	66302		RIKEN cDNA 2410005016 gene
<b>-</b>	10 BC003808			similar to hypothetical protein [Mus musculus]
<b>-</b> ;	10 NM 008548	17155	Man1a	mannosidase alpha man1a
<b>-</b> ;	0 NM 009493	22310	V2r4	vomeronasal 2, receptor, 4
F	10 AK004206	67282		AD16_HUMAN Protein AD-016 (Protein CGI-116) (x0009) 90 % /
= ;	10 AK015243	252876		hypothetical protein FLJ20125 [83% Homo sapiens]
= ;	10 NM 052975		,	
<b>≓</b> ;	0 NM_008036	14282	Fosb	nsp-like 1 protein nsp11 tma-sec and fosb; aa 1-338
¥ ;	10 AK017598			
¥ ∶	10 NM_021456			
5	10 NM_017463	18515	Pbx2	pre b-cell feukemia franscription factor 2 pbx2
5	010289 NM_010289	14610	Gja10	gap junction membrane channel protein alpha 10
5		15978	Ifng	interferon-gamma (muifn-gamma)
5	AK005661			
5	NM_009962	14764	Gpr44	putative g-protein coupled receptor crth2
1	NM_008078			
9	NM_009672	11737	Anp32	cerebellar leucine rich acidic nuclear protein lanp phosphoprotein 32 mpp32 encoded by genbank accession number u734; anp3;
10	NM_009004	19348	Rabekifl	rabkinesin-6
5	NM_028785	74146		RIKEN CDNA 1200017A24 gene
5		17751	Mt3	metallothionein 3
<b>1</b>	NM 011829	23917	Impdh1	inosine 5-phosphate dehydrogenase 1
10	BC005624			
10	10 NM 030717	80907	Lactb	serine beta lactamase-like protein lact-1
10	0 AK011064	70235		DKFZP434C245 protein [Homo sapiens] 90 %
5	10 NM_013415	11932	Atp1b2	atpase na+/k+ transporting beta 2 polypeptide
10	AK007097			
10	D AK003217			
5	0 NM_013459			
10	0 BC006600	60321	Wbp11	WW domain binding protein 11
2	0 AK013927			
2	10 NM_008508			
10	10 AF440737			
9	10 AK003496	73833		T08675 hypothetical protein DKFZp584F0522.1 - human (fragment) 49 %
=	AK011036	66230	Mrps28	mitochondrial ribosomal protein S28
11	AK016041			
	NM_015818	50785	Hs6st1	heparan sulfate 6-0-sulfotransferase 1
	AK009303			
11/	AK006173			
Ξ	NM_028717	74018	Als2	amyotrophic lateral sclerosis 2 (juvenile) homolog (human)
-	MM_010908			
	U69898	21856	4	translocase of inner mitochondrial membrane um44 nuclear encoding protein precursor form has size 50 Kg mature
	AY044265	235339		dihydrolipoamide S-acetylitansiensse (EZ component of pyruvate dehydrogenase complex)
4 1	NM_020611	57357	Srd5a2l	(eroid 5 aipha-reductase Z-like; H5AK gene; steroid 5 aipha-reductase Z like [Mus musculus] 100 %
	AKOOGAAG	69468		adult male testis riken crina clone:1700028b15
	2***	201100		BUGIL Hala Gasts Inval Cure Core

ne Description				p1 GPI-anchored membrane protein 1			_	_					hypothetical protein F23149_1(97% human)	_			RIKEN cDNA 4921513D11 gene	3b tumor necrosis factor (ligand) superfamily, member 13b			3 RAB, member of RAS oncogene family-like 3	and one brown relation for about a management of the sec		_	hypothetical protein FLJ22724 [Homo sapiens] 68 %		TATA box binding protein (Tbp)-associated factor, RNA polymerase I. A		Similar to hypothetical protein DKFZp434A1520.1 [70% Human]	cDNA 5730449L18 [Mus musculus]		<ul> <li>meiotic recombination 11 homolog A (S. cerevisiae)</li> </ul>			_	N-acety/glucosamine-1-phosphodiester alpha-N-acety/glucosamin/dase	•	frizzled homolog 4 drosophila fzd4	_		RIKEN cDNA 4921517J08 gene	
Gene		H2-M2	Nxph2	Gpiap1	Sirfada	Ŧ	Dunod	Cpne4		Rfx1	Miz1	Thra		Kcnj1	Rarg	Trim25		Tnfsf13b		Apcp6	Rabi3	AIA.	26.51	Cdc25c			Taf1a				:	Mre11a	Epb4, 113	Scy11	Galgt2	Nagpa	Çţ	Fzd4		Kane4		
Locus		14990	18232	53872	64452	15374	22242	74020		19724	17344	21833	54624	56379	19411	22660	70923	24099		74104	67657	10046	2500	12532	66722		21339		70348	66637	70775	17535	13823	78891	14422	27426	13043	14366	67711	57814	70891	
Cluster Access	11 NM_028333 11 NM_008896			11 NM_016739		11 NM 008258	11 BC012973	11 AK014396	11 AK005756	11 NM_009055	11 NM_008602	11 BC006025	11 NM_019458	11 NM_019659	11 NM_011244	11 D63902	11 AK014877	11 NM_033622	11 AK018430	11 NM_023732	11 AK016099	11 BC006253	11 AK009010	11 NM 009860	11 AK014865	11 NM_019718	11 NM_021466	11 NM_019453	11 AK011446			11 NM_018736	11 NM_013813	11 NM_023912	11 NM_008081	11 AF187073	11 NM_007803	11 NM_008055	11 NM_026330	11 NM_021342	11 AK016064	11 AK009780

Description solute carrier family 25 (mitochondrial carnitine/acylcarnitine translocase), member 20	choline kinase neurogenic differentiation 3 neurod3 RIKEN cDNA 2700023J09 gene chitinase 1 (chitotrisosidase) angiopoietin-like 3 RIKEN cDNA 2010004P11 gene sulfotransferase-related protein SULT-X1	hepatocyte nuclear factor hnf-1 RIKEN cDNA 170017111 gene CCR4-NOT transcription complex, subunit 4 Similar to CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) phosphatase, subunit 1, Isoform FCP [78% Human] galanin receptor 2 hypothetical protein HSPC 194 [Homo saplens] 85 % Mus musculus ubiquitinating enzyme E2-230 glut3 eccursor protein receptin 1 precursor protein mesenchyme fork head-1 protein; mfh-1	129/sv befa-n-acetylhexosaminidase hexb; beta-hexosaminidase beta subunit RIKEN cDNA 2310010M24 gene doublesex and mab-3 related transcription factor dmrt1 candidate sexual regulatory protein; transcript truncated SON protein [Mus musculus] 34 % 2-5 oligoadenylate synthetase 1G recoverin d-gtycosaminyl 3-o-sulfotransferase-3b 3-ost-3b kinesin 2 RIKEN cDNA 2410127L17 gene
Gene Sic25a20	Chk Neurod3 Chitt Angptl3 Sult-x1	Timp2 Tcf1 Cnot4 Galr2 Slc2a3 Cbin1 Foxc2	Hexb Dmrt1 Oas1g Rcvrn Hs3st3b Kns2
Locus 57279 76574	12660 18014 72569 71884 30924 67889 57429	21858 21405 66329 63221 67655 14428 66154 unknown 20527 12404	15212 71897 50796 66239 23960 19674 54710 16593 67383
Cluster Access 11 AK017913 11 AK021021 11 NM 020520 11 AK006096	11 NM 013490 12 NM_010396 12 AK012283 12 AK009012 12 AK002873 12 NM_026434 12 NM_020564 12 NM_020564 12 NM_020564	12 AK020699 12 NM_011594 12 NM_00327 12 AK003860 12 NM_016877 12 NM_010254 12 NM_010254 12 UM_010254 12 UM_01057 12 UM_01057 12 UM_01956	12 NM_010422 12 AK009282 12 NM_015826 12 NM_05233 12 NM_011221 12 NM_011221 12 NM_009038 12 NM_018805 12 NM_008450 12 AK010784



	oasic nelix-loop-ne	diappoint, Adol Tautiniy an Oct Tompies like windbi-intercent after from the Complex of the Comp	torsing amount in member 8	culin 2	RIKEN cDNA 9130204G15 gene	RIKEN cDNA A330104.06 gene	DNA segment, Chr 8, ERATO Doi 633, expressed	expressed in non-metastatic cells 2, protein	polymerase (DNA directed), lambda	adipose differentiation related protein	death-associated kinase 2	vav 3 oncogene	hypothetical protein FLJ20048 [Homo sapiens] 76 %	Eaf1 protein	transcription repressor crtr-1 developmentally regulated related to the cp2 family of factors		GL004 protein (Homo sapiens) 85 %	glypican 1	Similar to Rag C protein [94% Homo sapiens]	RIKEN CDNA 4921507P07 gene	Mus musculus adult male testis CDNA, RIKEN full-length enriched library, done:1700041N15 product:CHEMOKINE-LIKE FACTOR 2 VARIANT 2	nuclear receptor subfamily 4, group A, member 3	methylmatonic aciduria (cobalamin delicency) type B homotog (human)	gap function membrane channel protein epsilon 1	RuvB-like protein 1	RAB3B, member RAS oncogene family	origin of replication 3 homolog s. Cerevisiae orc3	lubby super-lamily protein fusp	Mus musculus adult male epididymis CDNA, RIKEN full-length enriched library, clone:9230110R08 product:nypothetical HiMG-1 and HIMG-7 UNA-binding domain (A+1-hoo	pleckstime and Sec7 domain protein 150% Homo saprens)	2.14 / MOUSE Line inger protein 14 / Lestragen responsive linger protein) (EIp) 3.2 %	SITUATION OF ASSISTANCE TRUSCULUS)	ninetan Livonokauta gene ninetan Livonokauta gene	Misconscience Software 3 and Misconscience Software 3 and Misconscience Software Sof	•		sialytiransferase 9 (CMP-NeuAc;lactosytoeramide althia-2,3-sialytiransferase)	testis expressed gene 13	hypothetkal protein FLJ13782 (94% human)	NADH dehydrogenase (ubiquinone) 1 beta subcomplex	hypothetical protein PTD004; homologous yeast-44.2 protein [99% Homo sapiens]	vascular Rab-GAP/TBC-containing: BUB2-like protein 1 [Mus musculus] 48.54 %	similar to glufathione peroxidase 2 [50% Homo sapiens]	oncogene-regulated cell adhesion molecule orcam		2119399A elongin B (Homo sapiens) 72 %	guidary-dearlyme a delydrogenase gon guidary-dearlyme a delydrogenase gon	Solected with the control of the con	RIKEN cDNA 1110025G12 gene	
_	20893 Bullo2		30934 Tor1b	71745 Cul2	77675	77785	52447	18103 Nme2	56626 Poll	11520 Adlp	13143 Dapk2	57257 Vav3	68732	74427 Eaf1	81879 Crtr1		75734	14733 Gpc1	75220	70821		18124 Nr4a3	77697 Mmab	118446 Gje1	56505 Ruvol1	69908 Rab3b	50793 Orc3	68842 Tulp4	77907	70965	71956	7967	19331	. S	56526 38961	Tac	20454 Siat9	83555 Tex13	252973 GhI2	66046 Ndufb5	67059	76795		57810 Cdon	57755 Dnajb7	72381	14488 Godh	70809 Octipa	77037	
Cluste Access	* NW 011496	1 AB012265	1 AJ297743	1 AK016520	1 AK020272	1 AK020734	1 AK017529	1 NM_008705	1 NM_020032	1 NM_007408	1 AF052942	1 NM 020505	1 AK004138	1 AK016628	1 NM_023755	1 AK020739	1 AK017226	1 NM_016696	1 AK015982	1 AK014840	1 AK006679	1 NM_015743	1 AK020286	1 NM 080450	1 NM_019685	Z NM 023537	Z NM_015824	2 NM 054040	2 AK020331	2 AK016466	2 AK010429	2 AKO06710	2 D26157	2 1.42336	2 NM 019942	3 NM 009311	3 NM 011375	3 NM_031381		3 NM 025316			3 AK009378			3 NIM 028218	3 NM 008097	3 NM_02/302	3 AK003912	

Clusto Access

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Mouse adult male kidney cDNA, RIKEN full-length enriched library, done:0610009A07 product:hypothetical Nitroreductase family containing protein
                                                                                                                                                                                                                                                                                                           multiple endocrine neoplasia type candidate protein number 18 dnaj famity protein mog 18; adult male small Intestine riken odna clone:2010301j22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WGAL_HIJAAN Neutrophil gelatinase-associated fipocatin precursor (NGAL) (25 kDa admis-2-inforoglobulin-related subunit of MMP-6) (Libocatin 2) (Oncogene 24p3) (32% human)
                                                                                                                                                                         dolichol-phosphate (beta-D) mannosyltransferase 1
112515 hypothetical protein DKFZp434B103.1 - (28% human)
BAK_MOUSE Bet-2 homologous antagonist/killer (Apoptosis regulator BAK)
Traf and Trif receptor associated protein; DNA segment, Chr 13, ERATO Dol 656, expressed [Mus muscul 100 % ubiquitin specific protease 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DD17_HUMAN Probable RNA-dependent helicase p72 (DEAD-box protein p72) (DEAD-box protein 17) 49 % /
                                                                                                                                                                                                                                                                                                                                                                                                                                               cysteine and histidine-rich domain (CHORD)-containing, zinc-binding protein t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    X-ray repair complementing defective repair in Chinese hamster celts 2 cDNA 2310008M10 [Mus musculus] RIKEN cDNA 1110021H02 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Niemann Pick type C2
RIKEN cDNA 9430034F23 gene
nudix (nucleoside diphosphate linked molety X)-type motif 7
replication protein-binding trans-activator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   solute camer family 17 (sodium phosphate), member 1
                                                             purinergic receptor P2X, ligand-gated ion channel, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similar to 148668 zinc finger protein 51 - mouse 43%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein MGC2827 [94% Homo sapiens]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     calcium/calmodulin-dependent serine protein kinase
                                         adult male thymus riken cdna clone:5830406p04
                                                                                                                                                                                                                                                                                                                                                                               9030607L17Rik RIKEN cDNA 9030607L17 gene
                                                                                                                                                                                                                                                                                                                                    Mpv17 transgene, kidney disease mutant-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          purinergic receptor P2Y, G-protein coupled 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 insulin-like growth factor binding protein-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mitochondrial ribosomal protein S18A spinocerebellar ataxla 2 hornolog (human)
                                                                                                                               piccolo (presynaptic cytomatrix protein)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      microsomal triglyceride transfer protein
                                                                                                                                                      cysteine-rich hydrophobic domain 1
                                                                                                                                                                                                                                                                                                                                                      RIKEN cDNA 3110005021 gene
                                                                                      feucine-rich repeat-containing 2
                                                                                                                                                                                                                                                                                                                                                                                                                          RIKEN cDNA 2810004A10 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        outer dense fiber of sperm talls 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RIKEN CDNA 2700007P21 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RIKEN cDNA 2300004C15 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RIKEN CDNA 2010001E11 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   vesicular membrain protein p24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SRY-box containing gene 3
                                                                                                           TATA box binding protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fyn binding protein fyb-130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RIKEN cDNA 0610042E07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fibroblast growth factor 1
                    odorant receptor S25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           YY1 transcription factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B lymphoid kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chloride channel 1
                                                                                                                                                                                                                                                                                          myosin vb
                                                                                                                                                                                                                    75393
56196 Ttrap
57775 Usp29
17919 Myo5b
57431 Dnajod
93734 Mpv171
77035
71564
66933
66917 Chordc1
                                       12482 Ms4a2
18436 P2x1
74249 Lrrc2
21374 Tbp
26875 Pclo
12212 Chic1
13480 Dpm1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22632 Yy1
68565 Mrps18a
20239 Sca2
Locus Gene
56861 Ors25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          83702 Akr1c6
14164 Fgf1
69459
68939
20504 Sic17a1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        212772
70839 P2ry12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23880 Fyb
16009 Igfbp3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20675 Sox3
67963 npc2
77289
67528 Nudt7
170742 RBT1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18286 Odf2
57434 Xroc2
66357
66155
12143 Blk
22360 Vmp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                140474 Muc4
12361 Cask
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17777 Mttp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12723 Clcn1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            67607
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                                                                                  1 AK010252
1 NM_013684
1 Y19185
1 Y11896
                                                                                                                                                                       NM_010072

AK014905

AK014905

AK012723

NM_021323

NM_021323

NM_021323

NM_021323

NM_021323

NM_021323

AK012666

AK012666

AK012666

NM_023634

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AK012666

AK012666

AK012666

AK012666
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NM_008343
AK012215
AK013804
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NM_020570
NM_025509
NM_025388
NM_007549
                                                           NM 00877
                    3 NM_02029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NM_009513
AK014820
                                           3 NM 00764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NM 030611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 NM 009198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 NM_026158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NM_010197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6 NM_009537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6 NM_026768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 NM 009125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AK020307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5 BC016456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5 BC008101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 AJ011106
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Description chemokine (C molif) receptor 1 breast cancer metastasis-suppressor 1 pheromone receptor V3R9 telomeric repeat binding factor 1 copz 1 nonclathrin coal protein zeta-cop lymphocyte artigen 6 complex, locus G RIKEN cDNA 2010012C09 gene	Mouse 13 days embryo head cDNA, RIKEN htth-length emrithed library, clone:31 10003A17 product:unknown EST COP9 (constitutive photomorphogenic) hornolog, subunit 3 (Arabidopsis thaliana) RIKEN cDNA 4933425L06 gene guanine nucleotide exchange factor (RCC1 retated) ubiquitin-conjugating enzyme E2G 2 ring finger protein 13 enhancer trap locus 1 distabless horneobox promylencytic floukemin	phospholipase C-like 2  T 17239 hypothetical protein DKFZp4348027.1 - human (fragment) 85 % similar to hypothetical protein FLJ12660 [90% Homo sapiens] RIKEN cDNA 18100 14F10 gene hypothetical protein FLJ12598 (12% human) - 138487 tastin - human 51.90 % DNA segment, Chr 3, ERATO Doi 250, expressed calclum channel, voltage-dependent, beta 2 subunit RIKEN cDNA 17000 19F09 gene prospero-related homeobox 1 WD capaci reformal 10	bably codiagen in alpha 4 chain; colda4  RNA binding motif protein 6  arachidonate lipoxygenase 3  serine/arginine-rich protein specific kinase 2  RIKEN cDNA 1810008021 gene tyrosine 3-monooxygenaselryptophan 5-monooxygenase activation protein, zeta polypeptide tripartile motif protein trim26 alpha phosphoprotein emiched in astrocyles 15  symptobovovi like 1  Symptopovovi like 1	Inter-alpha trypsin inhibitor, heavy chein 3.  Inter-alpha trypsin inhibitor, heavy chein 18.  Inter-alpha trypsin inhibitor, heavy chein
Locus Gene 23832 Kcri 107392 Brms1 81010 V3R9 21749 Terf1 56447 Copz1 17072 Ly69 68440	73112 26572 Cops3 66763 66763 22213 Ube2g2 24017 Rnf13 13390 Ell1 13390 Dix1 18854 Pml	224860 Pid2 76454 76454 76826 69064 68927 78733 52184 12296 Cacub2 71860 19130 Prox 1	12829 Codas 19654 Rbm6 23801 Aloxe3 20817 Srpk2 69094 22831 Ywhaz 22870 Trim26 18611 Pea15 20655 Sybi1	Lrp 1b
Cluste Access 6 NM_011788 6 AF23380 6 NM_030735 6 NM_099352 6 NM_019817 6 NM_011660 6 X70920 6 BC005775	7 AK013984 7 NM_011991 7 NM_025751 7 NM_019803 7 NM_019803 7 NM_011883 7 X69942 7 NM_010053 7 NM_010053 7 NM_010053	7 AR013615 7 AR010827 7 AK010827 7 AK010827 7 AK01494 7 AK01494 7 L20343 7 AK006118 7 NM_008937 7 AF296775	7 235167 7 NM_011251 7 NM_011786 7 U92456 7 AK007386 7 NM_011740 7 AF230395 7 NM_006556 7 AF230395	5 0-68

Description

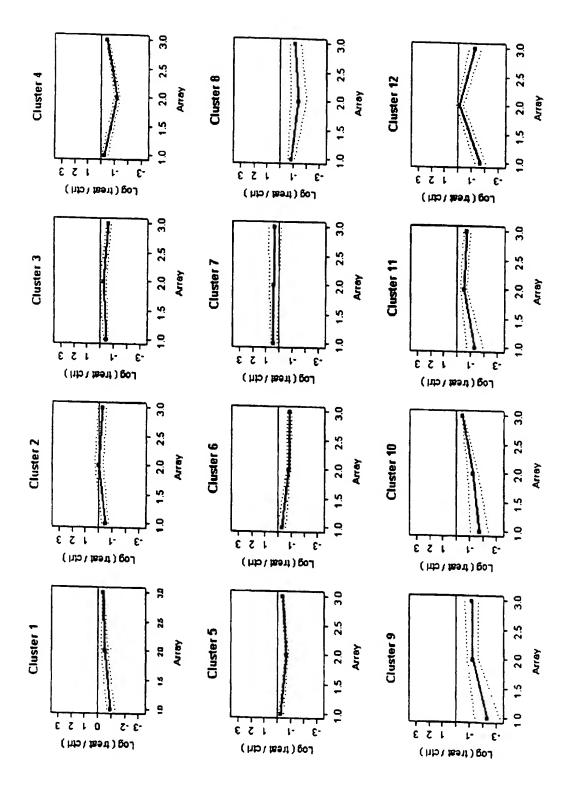
Cluste Access

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PGPI_HUMAN Probable pymolidone-carboxylate peptidase (5-oxoproby-peptidase) (Pyrogutarny-peptidas 95 % keratin associated protein 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             growth differentiation factor-9b gdf-9b; bone morphogenatic protein 15 bmp15
                                                                                           Ngf-A binding protein 1
RIKEN cDNA 5730409E15 gene
amyloid beta (A4) precursor protein-binding, family A, member 3
                                                                                                                                                 ring finger protein 17
B-cell leukemia/lymphoma 6
sp:P46096 - SY71_MOUSE Synaptotagmin I (Sytl) (p65) 38 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        complement component 1, q subcomponent, gamma polypeptide HSPC038 protein [Homo sapiens] 100 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similar to hypothetical protein FLJ20546 [82% Homo saplens]
                                                                                                                                                                                                                                                                                                                                        similar to keratin associated protein 4.7 [31% Homo sapiens]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     putative seven transmembrane apanning receptor puma-g insulin-like growth factor i receptor igf4
                                                                                                                                                                                                                                                                                                                     CA00_HUMAN Protein CGI-100 precursor (89% human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  serine/threonine kinase FKSG81 [Homo sapiens] 44 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hypothetical protein MGC15435 [Homo sapiens] 51 %
                                                                                                                                                                                                                                                                                                                                                                            achaete-scute complex homolog-like 2 (Drosophila)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rab6-interacting protein 2 [Mus musculus] 23.36 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protactin-like protein M
hypothetical protein MGC14827 (85% human)
RIKEN cDNA 2410012M07 gene
                                                                                                                                                                                                                                                                                                                                                                                                               perinatal skeletal myosin heavy chain 3 end
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mad homolog 3 drosophila madh3; msmad3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ubiquitin specific protease 5 (isopeptidase T)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytotoxic T-lymphocyte-associated protein 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    zinc finger protein 341 [Homo saplens] 90%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JC6547 high sulfur protein B2E - rat 37 %
                                                                                                                                                                                                                                                                                                     transcriptional regulator, SIN3B (yeast)
                                                                                                                                                                                                                         F-box and WD-40 domain protein 5
                                                                                                                                                                                                                                                                                                                                                                                               bone marrow stromal cell antigen 1
                                                          Dullard homolog (Xenopus laevis)
                    glucagon-like peptide 1 receptor
                                    RIKEN cDNA 4930519F16 gene
                                                                                                                                                                                                                                                                                  RIKEN cDNA 4930443G12 gene
                                                                                                                                                                                                                                            unc-84 homolog A (C. elegans)
   tripartite motif protein trim 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                glia maturation factor, gamma
                                                                                                                                                                                                                                                                                                                                                                                                                                  leukotriene A4 hydrolase
heat shock protein 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              vaccinia related kinase 2
                                                                                                                                                                                                                                                                                                                                                             olfactory receptor 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                major vault protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                androgen recept
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGI-94 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         reticulocalbin 2
                                                                                                                                                                                                                                                               CD9 antigen
                                                                                                                                                                                                          matrilin 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   caspase 9
                                                         67181 Dullard
                                                                                                                                                                                                                                                                                                                                                                                                           17885 Myn8
16993 Lladh
16512 Hspa2
56635 Pripm
68201
71979
74239
74239
77127 Madh3
67205 Cgi94
12477 Ctlad
26618 Rcn2
66628
72114
66522
16699 Krtap13
 94091 Trim11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22225 Usp5
80885 Puma-g
16001 lgf1r
                                                                                                                               57267 Apba3
30054 Rnf17
                                                                                                                                                                    12053 Bcl6
77314
17183 Matn4
30839 Fbxw5
                                                                                           17936 Nab1
70502
                                                                                                                                                                                                                                              7053 Unc84a
                    4652 Glp1r
                                                                                                                                                                                                                                                                                                                                                           27216 Olfr154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16001 lgf1r
12155 Bmp15
78388 Mvp
                                                                                                                                                                                                                                                              12527 Cd9
74686
                                                                                                                                                                                                                                                                                                  20467 Sin3b
                                                                                                                                                                                                                                                                                                                                                                          17173 Ascl2
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69696
12262 C14g
68036
69922 Vrk2
63986 Gmfg
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8 NM_053168
8 NM_021332
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8 AK017530
8 NM_018758
8 AF285585
6 NM_009744
8 AK021056
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NM_013908
AF343752
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8 NM_008763
8 M12289
8 NM_008517
9 NM_018991
9 NM_018991
9 NM_026613
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9 NM_013476
9 NM_013476
9 NM_026031
9 NM_00843
9 NM_011992
9 AK010876
9 NM_013217
9 NM_013671
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9 NM 030701
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9 AK016890
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9 NM_080638
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9 AK012664
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DYNC_Human Dynactin complex 50 kDa subunit (50 kDa dynein-associated polypeptide) (Dynamilin) (DCTN- 94 %
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similar to hypothetical protein MGC10999 [Homo saplens] 81.25 %
- A55190 transitional endopdasmic reticulum ATPase (EC 3.6.1.-) [validated] - rat 35 %
hypothetical protein FLJ12783 [Homo sapiens] 96 %
                                                                                                                                                                                                                   Similar to WD domain, G-beta repeat-containing protein [Homo saplens] 83 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    71709 similar to A49307 98K GTPase-activating protein ABR, brain - human 29 % 58804 Cdc42ep5 CDC42 effector protein (Rho GTPase binding) 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical protein CLONE24922; hypthetical protein [30% Homo saplens]
                                                                                                                                                                                                                                                                                                                                                                         calsequestrin 2
Similar to APXL_HUMAN Apical-like protein (APXL protein) humna 28 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Huntingtin interacting protein K; hypothetical protein [Homo sapiens] 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  zinc finger 202 mt znf202 scan-krab-zinc protein znf202-m1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         likely ortholog of kinesin light chain 2 [69% Homo sapiens]
                    11517 Adcyap1r1 adenylate cyclase activating polypeptide 1 receptor 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ARP6 actin-related protein 6 homolog (yeast) regulating synaptic membrane exocytosis 2 hypothetical protein FLJ22353 (Homo sapiens) 85 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cadherin egi lag seven-pass g-type receptor celsr t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     thymidylate kinase family LPS-inducible member
                                                                                  interferon regulatory factor-3 irf3 irf-3; factor 3
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                                                                                                                                                                                                                                          transitional epithelia response protein
                                                                                                                                                                                                                                                                                  butyrophilin, subfamily 1, member A1
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                                                                Nil protein 2 (89% Homo saplens)
                                                                                                                                                                                                                                                                                                                        RIKEN cDNA 2010107K23 gene
                                                                                                                                                                                                                                                                                                                                                                                                                 neuregulin 4 nrg4
RIKEN cDNA 6330583M11 gene
RIKEN cDNA 1110068E08 gene
RIKEN cDNA 1700006J14 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  T-cell receptor beta, variable V20
                                                                                                                                                                                             Tetraspan NET-6 (95% human)
                                                                                                                             Nuclear neuronal protein 15.8 interleukin-10 receptor alpha
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54131 lrf3
18575 Pde1c
104130 Np15
16154 ll10ra
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71707 Tere1
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23864 Odz2
72043 Sult2
80902 Zip202
12425 Cckar
7478
77478
77252
72962 Eggf1
14862 Gstm1
14862 Gstm1
14862 Gstm1
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81500 Sil1
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67019 Actr6
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12614 Celsr1
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9 NM_023175
9 NM_016849
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9 NM_01935
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10 NM_007986
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10 NM_00550
10 NM_00814
10 AK004934
10 NM_0224465
10 AK004654
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10 NM_021454
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11 NM_009827
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11 AK020480
Cluste Access
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Description RIKEN cDNA 2310016G11 gene Similar to IS8401 protein-tyrosine kinase (EC 2.7.1.112) JAK3 - mouse 69% olfactory receptor 17 similar to contrapsin-like protease inhibitor related protein (CPt-23) [Rattus norvegicus] RIKEN CDNA 2310003M01 gene RNA birding motif protein 3 RIKEN cDNA 4930405H06 gene		THIO_HUMAN Thioredoxin (ATL-derived factor) (ADF) (Surface associated sulphydryl protein) (SASP) 52 % zinc finger protein 120 wist gene homolog 1 (Drosophila) wist gene homolog 2 (Drosophila) wist gene homolog 3 (Drosophila) wide are factor of kappe light polypoptide gene enhancer in B-cells inhibitor-like 1 protease (prosome, macropain) 288 subunit, ATPase 1 2-cell-stage, variable group, member 1; variable group of 2-cell-stage gene family HS1 binding pro RIKEN cDNA 1500035H01 gene upregulated in colorectal cancer gene 1 parvin, alpha	BCL2-tike 12 (proline rich); Bcl-2 like proline-rich protein 12 [Homo sapiens] 81 % serine (or cystein) proteins as inhibitor, clade E, member 2 small gutamine-rich tetratroopeptide repeat 2 small gutamine-rich tetratroopeptide repeat 1 hip 3 small gutamine-rich tetratroopeptide repeat 3 small gutamine-rich tetratroopeptide repeat 4 block-benteracting protein 1, NIP3 atp synthase the transporting mitochondrial f1f0 complex subunit e atp5k; ffm-1 f1f0-atpase furnarate hydratase 1 programmend cell death 10 plakophilin 1 straided muscle-specific sertne/threonine protein kinase speg ASSSTS ankyrin 3, fong splice form - (28,61% human)
Locus Gene 69578 76799 18314 Olfr17 71886 19652 Rbm3 73818	108660 77632 74589 78783 Brpf1 69769 12951 Crx 74720 52613 Rps6ka4 74420	67402 104348 Zip120 22160 Twist! 19038 Nikbi! 19179 Psrmc! 23897 Hs1bp! 76568 94228 Ucc! 57342 Parva	75736 20720 Serpine2 52551 ggt 12176 Bnip3 11958 Atp5k 14194 Fh1 56426 Pdcd10 11790 Apeg1 73845
Cluste Access 11 AK009387 11 AK016715 11 NM_020598 11 NM_028740 11 NM_018097 11 NM_016809 11 AK015098	11 AB030190 11 AK019795 11 AK014702 11 AK014230 11 AK015522 11 AK015527 11 NM_078484 11 NM_078484 11 NM_078924	11 NM_026132 11 NM_02326 11 NM_01658 11 NM_010909 11 NM_008947 11 AF067063 11 NM_01826 11 NM_023931 11 NM_023931	12 AK017362 12 NM_009255 12 BC00336 12 NM_007507 12 NM_007507 12 BC006048 12 NM_019645 12 AF215896 12 AK007013 12 AK016847





Chater Access	Locus Gane	Description
1 AK015845		
NM_008664	17930 Myom2	myomesin 2 myom2
1 AJ242625	13406 Dmp1	denlin matrix prolein 1
1 NM_007975	14065 F2rt3	protease-activated receptor 4 pardem-coupled receptor thrombin
1 NM_017468	13801 Enam	enametin - ENAM_MOUSE Enametin precursor 100 %
1 NM_017404	27393 Mrpl39	10 day old male pancreas riken caha done:1810033d11; unknown c21orf8
1 X76011		
1 AK004006		
1 NM_018774		
1 NM 008976	19250 Ptpn14	outsive protein brossite phosphalase oto38 a sh3-binding site in the spacer region connecting reterminal hand 4.1-lite domain and elements in
1 AK014894		
1 NM_016778		
1 AK007277	73816	RIKEN CDNA 1700125F08 nene
1 NM 008372	16197 1174	interior recent annual and in 172 - 172
NM OTODO	13090 Cun2e40	Anthonic Dept. 2 subfamily a cohemica 40
1 AKO 10834	7677E	LYCATION AT COLUMN 1, SOUGHING III, POSPERIOR TO
75000 FEB 9	200	MINERA COUNTY 24 TO 1950-OZ BENE
1 NM 01303/		
A-0011706	12/23 ClCn1	chloride channel 1
NM_013714	26573 treb/1	interferon response element briding (ador 1
1 AK016979		
1 NM_023055	65962 Slc9a3r2	solute carrier tamity 9 isoform 3 regulator 2, isoform A; sodium/hydrogen exchanger [Mus musculus] 100 % /
1 AK005213	66239	Iruncaled SON protein (Mus musculus) 34 %
1 AK009119	69473	kerain associated protein 3.1 (87% Homo sapiens)
1 NM 032400	84112 Gar91	G arotein-coupled recentor 91
1 A.750690		New restriction to a to the second of the se
1 BC004774	07009 075103	an account of the second secon
1 114 007500	2012 10 10 10 10	
1 X74366	MILLIO 2/07	crownergo: receptor mascanine a curria
00777	0000	
8/2020 WA	20030 303970	Small inductore cytomine aco scyazo
AKOIOLIS		
I NM 009352	21749 Tert1	teloment repeal binding lactor 1
1 AK00/894		
I NIM DIDAYS	15464 Hrc	histidine non calcium binding protein
1 AK016893	71166	RIKEN cDNA 4933424G06 pene
1 NM_019455	54486 Ptgds2	prostaglandin D2 synthase 2, hematopoletic
1 NM 008506	16918 Lmvc1	SET
1 NW 013614	18263 Odc	omitibie decendoxytase, structural
1 NM OOR320	16192 11500	Interfaction, Economic concepts
1 MM 026572	68133	and the matter site of the desired to the second se
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1 AK014327	2446/2	RIKEN CONA 323MOTLUS pene
1 AF282302	258434 MOR224-6	oliaciay recepior MOK224-6
1 NM_010155	13875 Erl	esiz repressor lactor en
1 NM_010406	15139 Hc	complement component c5s pro-c5 precursor
1 AK006388		
1 AK016424	20803	RIKEN CDNA 4931402H11 gene
1 BC006060	107049	hypothetical protein FLJ10595 (86% human)
1 AK020377		
1 NM_031373	72075 Ogfr	opioid growth factor receptor
1 NM_026187		
1 NM 024468	79263 Trim39	ripantie modif protein 39
1 NM 020588	57439	RIKEN cONA 1300007812 gene
1 NM 024210	67862	RIKEN cDNA 2310033P09 [Mus muscudus] 100 %
1 AK008209		RIKEN CONA 2010012G 17 pene
1 AK010336		
1 NM 009476	22269 Unk2	uroclakin ii muož
1 AK015351		adult male testis riken considerations.
1 NM 019873	Fither	EX508 binding professional and a second seco
1 AKO03611		
1 AK002744		

Locus Gene Description	developmental pluripotency associated 2	Smell Cut	S8996 hypothetical protein, MNCb-1301 (100% Mus musculus) 22376 Was Wiskell-Aldrich syndrome bonneton frimman				210336		30949 Lcmii feudine carboxyl metrytiransferase izmti	64178 Gard8 - Carrelain restricted teacontre 88	_			68/35 KIKEN CDNA 2810039M1/ gene	17850 Mui methytmatonyl-coenzyme a mut	B3galt2	75858 RIKEN CDNA 4830568L21 gene		Cdkn2b	70360 hypothetical protein d.M.73B4 (95% Homo sapiens)	107589 Mytk myosin, light pohypeptide kinase	11998 Avp arginine vasopressin		20248 Servinty serine (or cysteine) proteinase inhibitor, clade 8. member 4	Fros	Cecngi	\$ 20191 Syda saroopyran ayina (ryanopina ayina goronopina ayina goronopina ayina goronopina ayina goronopina ayina ayina goronopina ayina goronopina ayina ayi	Nel Nel		æ		54651 USPZ/X ubiquiln specific protease ZY, X diramasame	Mus muscutus adult male testis cDNA, RIXEN full-length emiched library, cone;4930528A17 product;unknown EST			<u>2</u>	25049 Dmrtz doublesek and meta-7-fested trenscription factor 2	757.0 KIARN GUNA 4855A24KI 5 gene	15976 Ithar 2 mierteran alpha and beta receptor 2 lihar2: type i ithar2c	11409 Acads acety-coenzyme a dehydrogenase short chain acads
Chrster Access	1 NM_033572 1 AK010743	1 AF 109905	1 AK016967 1 NW 009515	1 NM_009504	1 AK007700	1 NM_025504	1 AK018470	1 NM_008668	7 NM_025304	2 NM 022427	2 NM 011225	2 AK009653	2 NM 020493	2 AK005168	2 NM 008650	2 NM_020025	2 AK016255	2 NM 015830	2 NM_007670	2 BC016329 2 NM 013556		NM_009732	NM_010569	NM 009126	NM_011729	NM_007582	NM COSTS	NM 023126	NM 011161		NM 015785	2 AF 229643			 -	•	•	X81632		

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Rabb-interacting protein 2 [Mus muscutus] 24 %
RIKEN CDNA 4830541(104 gene
Mus muscutus aduli male testis cDNA, RIKEN full-tength enriched library, clone:4930550C14 product.hypothetical 1Q calmodulin-binding matif containing pratein
H2B histone family, member C [Hanno sappens] 44 %
Mus muscutus aduli male hippocampus cDNA, RIKEN full-tength enriched library, clone:2900034J12
Mus muscutus aduli male hippocampus cDNA, RIKEN full-tength enriched library, clone:2900034J12
histone deacelylase hdac?
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KIAA0547 gene product [80% Homo septens]
leucher-rich and death dornain containing; p53 protein induced, with death domain [Mus musculus] 40.00 % todd Ozbren-in harmolog 1 (Drosophia)
add Ozbren-in harmolog 1 (Drosophia)
radii dehydrogenase udvetunone (es protein 4 18 kda ndu54; 13 days embryo liver riken cdna chone;2510049c12
CHD: MOUSE CHROMODOMAIN-HELICASE-DNA-BINDING PROTEIN 1 (CHD:) 40 %
                                                                                                                                                                                                                                                                                                                                                  Similar to HUMAN Siress-induced-phosphoprotein 1 (ST11) (Hsp70/Hsp90-organizing protein) [44% Human] stip!
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cyclin-dependent kinasa inhibitor 18 (P27)
nuclear transcription factor y gamma nfyc; factor yc
CD163 antigen
brain protein 16; DNA segment, Chr 15, ERATO Do 741, expressed [Abs musculus] 100 %
                                                                                                                                                                                                                  polymerase della interacting protein 38 gamma subunit of sodium potassium alpase adulf lemale placenta riken coha done: 1600023b02; small inducible cytokine a27 scya27 RIKEN cDNA 493051 IH01 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypothetical protein, MGC:7623; hypothetical protein MGC7623 [100% Mus musculus]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mammalian atonal homologue 4a homolog helix-loop-helix protein; neurogenin ngn2
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SMC2 structural maintenance of chromosomes 2-tite 1 homotog (yeast)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ma polymerase il 4 14 kda subunit rpo2-4
T34522 hypothetical protein DKF2p566D244.1 (99% human)
                                                                                                                                                                                                                                                                                                                                                                                                                   chemokine (C-X-C motif) ligand 10 terminal modulator protein [Homo saplens] 39 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          acety-Cocnzyme A synthetase 2 (ADP forming)
SH3-domain GRB2-like 1
                                                                                                                                                                                                          ectin, galactose binding, soluble 12
Description
hypothetical protein MGC7434
mesoderm posterior 2 mesp2
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                                                                                                                    calbindin 2
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67611 Pdip36
11936 Fxyd2
20301 Scya27
75820
66326
69545
57816 Tesc
Locus Gene
211556
17293 Mesp2
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20135 Rrm2
30826 Pcdh13
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67287
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18046 Nfyc
93671 Cd163
59053 Brp16
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71519 Cyp2uf
14211 Smc211
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20405 Sh3gi1
75894
                                                                                                               12308 Calb2
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Description ring finger protein (C3HC4 type) 19	ubiquilin C galanin receptor 3 neighbor of a-kinase anchoring protein 95 nakap95; 10 days embryo riken cdna done:2610005122 neighbor of a-kinase anchoring protein 95 nakap95; 10 days embryo riken cdna done:2610005122	Moloney sarcona oncogene adult male testis riken odna clone:1700007k08	inddearnine-pyrrole 2,3 dioxygenese enand incuded 2 (putative ribonuclease III; putative protein p241 which interacts with transcription factor Sp1 [Hom 99 % I enand incuded 2 (putative ribonuclease III; putative protein p241 which interacts with transcription factor Sp1 [Hom 99 % I	N-methytpurine-ONA glycosylase lyrosine 3-monooxygenaselhytiophan 5-monooxygenase activation protein, zeta polypeplide	13 days embryo head riken cdna clone:3110056m11 selenoprotein P. plasma. 1 sip-binding cassette sub-family 1 gcn20 member 2 done mgc.7932; protein abcf2	RIKEN cDNA 2410019A14 gene KIAA1001 protein [Homo sapiens] 78 % RIKEN CDNA 1500031J01 gene microsphende protein done mgc:5852; nucleater msp58	ATP synthase. H+ transporting, mitochondrial F0 complex, subunit f, isoform 2 copine till recompatibility 2 o region beta locus h2-ob histocompatibility 2 o region beta locus h2-ob 14 syst 2 hypothetical protein F5K20.320 -(23%, Arabidopsis thatlana) 14 syst 2 hypothetical protein F5K20.320 -(23%, Arabidopsis thatlana) polt_ akouse Retrovinus-related POL polyprotein [Contains: Reverse transcriptase ; Endonucleaself 32 % M.musoulus) etoposide induced 2.4 mRNA	atp-binding cassette sub-family a abc 1 member 2 abca2  GA repeat binding protein, beta 2, pir/A53950 - A53950 transcription factor GABP beta 2-1 chain - mouse 100.00 %  GA repeat binding protein, beta 2, pir/A53950 - A53950 transcription factor GABP beta 2-1 chain - mouse 100.00 %  RINEN DIVA 433405012 gene caspase 3, apportors related cysteme protease caspase 3, apportors related cysteme protease ATP synthase. N+ transporting, mitochondrial F0 corruptex, subunit c (subunit 8), isoform 2	adull male testis riken odna done:493340 tb01  RNA binding motif protein 16  zinc finger protein 146 zfb146  zinc finger protein 146 zfb146  micromotar calcum activated neutral protease large subunit camp1 protease mu-calpain; calpain I capn1  micromotar calcum activated neutral protease large subunit camp1 adaptor protein complex ap-1 beta subunit done mac:5850; ap1b1
Locus Gene 30945 Rnf19	22190 Ubc 14429 Gafr3 54194 Nakap95	17451 Mos 19119 Pmn2	15930 Indo 14000 Etahi2	17477 Mpg 22631 Ywhaz	1977 1 Rlbp1 20363 Sepp1 27407 Abcf2	69746 74008 67529 51812 Mers1	57423 Atp5j2 70568 Cpne3 15002 H2-Ob 67530 70770	11305 Abca2 213054 Gabbb2 230991 12367 (Gabb) 67942 Alp592	71027 106583 Rbm16 26465 Zfp146 12333 Capn1 11764 Aprib1
Chuster Access 3 NM_013923	3 AK003581 3 NM_019639 3 NM_015738 3 NM_017476	3 AK016522 3 NM_020021 3 NM_008933	3 AJ223472 3 NM_008324 3 AK003651 3 NM_010027	3 NM_025790 3 AK016431 3 BC002245 3 NM_010822 3 BC016080	3 AMC07580 3 NM_020599 3 NM_009155 3 NM_009431 3 AF213382	3 BC004728 3 AK010555 3 AK003511 3 AK018132 3 NM_026218 3 NM_016766	3 AR017758 3 NM_026526 3 NM_020623 3 AR017357 3 AR0166310 3 AR016634 3 AR016634 3 AR016634	3 AK©A706 3 AKO0847 3 AKO08847 3 AKO0878 3 AKO04079 3 AKO10289 3 AKO10289 3 AKO10289 3 AKO1689 3 AKO1689 3 AKO1689 3 AKO1689 3 AKO1689	3 AK014265 3 AK013485 3 AK011348 3 NM_011160 3 BC002302 3 NM_011980 3 BC002301 3 NM_071800 3 NM_07500

Control Assessed	Locus	200	Description
2 5050 020585	57437		HSPC041 protein (95% north-september 200 done: 4933432e16
3 AVON6727	67003		adult male testis riken consideration on the first of the protein
3 KM 019752	64704	64704 Prss25	serine protease triads escretaria como como como como como como como com
3 NM 010757	17135 Mafk	Maft	v-maj musculoaporeur en marcon
3 AF071068			
3 AKUT1363	14433	14433 Gepd	gyceraidehyde-3-phosphale denydrogen was
3 NM 021386	58187	58187 Cldn10	claudin-10
3 AK021021			
3 80010810			in the section of the
3 NM 009826	12421	12421 Cc1	cc1 coiled-coil for for in under the market coiled-coiled coiled-
3 NM 011750	22668	12668 Zfp162	zinc finger for our in expressed developmentally down-regulated nedd8
3 NM 008683	16002	18002 Nedd8	neural precures testinated
3 NM 009736	1201	12017 Bag1	BCL-SSCARrate are cleaving protease (81% Homo saptens)
3 NM_023220	66552		Jacks and Artifacts and an analysis of the second and a s
3 NM 007458	1171	11771 Ap2a1	BIDITAL CONTROL OF THE CONTROL OF TH
3 NM 025982	,		RIKEN CDNA 2410003H12 gene
3 AK010358	160		RIKEN cDNA 5730493819 [Mus musculus] boxus in contract co
4 AK00/55/	2087	Eners Tund3	ubiquitous tropomodulin u-fmod actin litarines pomice con a construction to the contract of th
4 NM_016963	15150	15159 Hccs	Holocylochrome c synthetase
4 AF277718	101502	101502 Hsd3b7	hydroxy-della-5-sterior defluydy-usa-s.
4 AK017113	66748	_	RIKEN CONA ASSOCIATION OF THE PROPERTY OF THE
4 AK019886			homeofrom domain interacting protein
4 NM 031879	8394	83946 Phip	precisin included in muscle
4 NM 009948	12895	S Coilb	Carmine political interpolity (Drosophile)
4 AK017858	13831	Epc	erra MOUSE Serine protease inhibitor Kazal-type 4 precursor (* 1970)
4 Y11505	20731	Spinke	ENIKEN CONA 24 10030K01 gene
4 AK011728	67.623	,	email chemokine (C.C. motif) ligand 11
4 NM 011330	2029	20292 Corre	calcium channel, voltage-dependent, alpha ti suoun
4 AY026384			
4 AK008468			
4 NM U26300	74694	•	T48686 hypothetical protein Livraginal Advantage 3
4 AROIS/00	26.43	26433 Plod3	procedegen-lysine, 2-dyogularia
4 NIN D1106	67358		RIKEN CDNA 170009377   100 ms
4 MIN ON 1857	2396	23965 Odz3	odd ozhen-m homosg s drosymed; islet specific glacose-8-phosphatase (mus inserned)
4 NM 021331	14378	8 Gepc-rs	ghoose-6-phosphalase causing in the case of the case o
4 MM 009020	1937	19374 Rag2	recombination activated by Vernal
4 NW 010084	1352	13524 Adam18	disintegrin and metallicity of the control of the c
4 AK016580	11078	110782 Aldh5a1	addit male fests there was additional additi
4 AK014380	74016	4	11/200 nytotrence in the state of the state
4 NM 019566	562	56212 Arbg	Appropries
4 AK002703			
4 NM 008059			
4 NM 028119	9000	1 Moent	mesodern development candidate a sociation and an analysis by the form of the
4 NM 030705	1337	13370 Dio 1	adult male kidney riken odna clone;00 tuu rike; usaan aan aan aan aan aan aan aan aan aan
4 NM 00/800	3		•
4 AKODASIB	123	12363 Casott	caspase 11 casp11; caspase 11
4 NEW 00/903			Administration of the second of the second s
A NM 007544	121	12122 Bid	BH3 mercentag control
4 NM 009412	219	21985 Tpd52	lumor protein Los- Lista sectional lengthe zipper, extra-nuclear factor
4 NM 031378	25	83492 MIZE	TICHER COLLEGE
4 NM 010062			
4 AK010807			
4 AMUIO400	58	58181 #20	interleukin 20 UZO
5 AK011591	76964	3	RIKEN CONA 2610020124 BUT
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Cluster Access 5 AK018457	Locus Gene	Description	
5 AF25125	64452 Stc5a4a	4a solute carrier family 5, member 4a	
5 AK014709 5 AK007941			
5 NM_019849			
5 NM_026177 5 NM_019519	67467	RIKEN cONA 1200011118 gene	
5 KM 025459			
5 NM_007480	11844 ArfS	adp-ribosylation factor 5 arf5	
5 AF217002			
5 NM_010811	17423 Nds12	glucosaminyl n-deacetylase / n-suffo	glucosaminyi n-deocetylase / n-sulfotransferase dual enzyme activities; heparan sulfate n-deacetylase/n-sulfotransferase
5 NM_010268			
5 NM_021325	20479 Vrzedb		antigen identified by monoclonal antibody MRC OX.2 receptor Antigen identified place are search to receive and search and to the search and the
5 BC002148	1		oriente canon no emistrate accessor mante provococop pe asser y pas, vacuota sorieny en yeast date migroor a
5 NM 011986			
S NM_025591	66488	RIKEN cDNA 2010309E21 gene	
5 NM_030251	80283 Abib1	ankyrin repeat and 8T8 (POZ) doma	ain containing t
5 NM 020508			
5 NM_008737	18186 Nrp	neuropilin	
5 AK013041	69940	Sec3-like; homolog of yeas! exocyst (	Sec3-ike; homolog of yeast exocyst protein Sec2p (Homo sapiens) 86 %
5 AK017789	1000	Calif. Ulitable Drobable conference	CANT LIMBAR Describes controlled controlled blacks CANT (I.M. ISEN Income)
5 NM 008217	15118 Has3	hyaturonan synthase 3 has3	ליווים לינונייו אוופספ כאיר ובת (כל א וערופון)
S NM_033077			
5 NM_008556	18611 Pea15	phosphoprotein enriched in astrocytes 15	52 82
5 NM 028732	74048	RIKEN CDNA 4632428N05 gene	
5 AF090691	13012 Cst8	cystatin 8 (cystatin-related epididymal spermatogenic)	si spermatogenic)
5 X98456	0.00	M.musculus ORF1 and ORF2 genes	
5 AK011480	14538 GCM2	gucosaminytransterase, t-orangmag enzyme	ยาวุราช
5 NM 008260	15377 Foxe3	(orthead box a3 foxa3	
5 AK011565	69904	RIKEN cDNA 2610027F03 gene	
5 AK018093			
5 NM 025295			
5 BC017127			
5 AK005930			
S MAN DOOR	01010	C) military designation of the second	As and and a second as a secon
5 NM 009539			915
5 NM 018764	54216 Pcdh7	protocadherin podh7	
5 NM_021537			
5 NM_021452	58802 Kcnmb4	polassium large conductance calcium	polassium lange conductance calcium-activated channel subfamily beta member kombb
5 NM 010060	13411 Deabet1	the first the same of the same	
5 AK005564			
6 NM_011128			
6 NM 022014			
6 NM 025865			!
6 NM 026518	15551 Hirtib	5-hydroxytryptamene (serotomn) receptor 18	MOT 18
6 AK003792		•	
6 NM 019445	20604 Sms1 54418 Fmn2	preprosomatostaun formin 2	
6 AK020538	7397	Lysozyme C. type M precursor (1.4-bel	Lysozyme C, type M precursor (1.4-beta-N-acetymuramidase C) (62% Mus musculus)
6 AF057287			
6 NM_030706	80890 Trim2	Inpartite motif pratein trim2	

Description

Locus Gene

Chuster Access

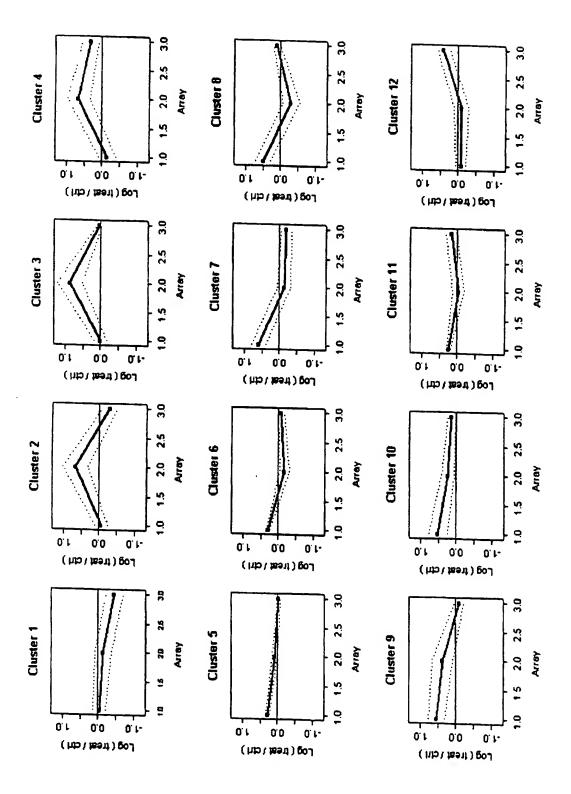
Description  I transforming growth factor beta 1 induced transcript 4  arr grotein putainva alpase and helicase  Mus musculus 13 days embyro head cDNA, RIKEN full-length enriched library, clone:3110043021 product:unknown EST  Mus musculus gene for Cox 17p, complete cds	S-adenosyhornocystehre hydrolase 1 a txt cotagen chain 2 similar to Ran bridring protein 11 (IPO11) [95% Homo saptens) phycoprotein-associated amino acid transporter lat2	TATA box binding protein (Tbp)-essociated factor, RNA polymerase I, C RIKEN cDNA 1110034G24 gene	fibroblast growth factor 22 preproadrenomedulin; adrenomedulin adm	balb/c collagen iv alpha 4 chain; co4a4 RNA binding moill protein 6 midline 1 syntrophin, actdic 1		1 ATPase, H+ transporting, V1 subunit C, isotomn 1 TEC formal family, member 8 TEC formal family, member 8 Technical parallel family, member 100 % Technical family, member 100 Technical family famil	RIKEN cDNA 4821513E08 (Mus muscutus) 100 % endoptasmic reticulum (ER) to nucleus signalling garman-gutamy carboxylase garman-gutamy carboxylase reticulum (ER) to nucleus signalling reticulum (ER) to nucleus signalling (ER) to nucleus sig	2004395A metanin-concentrating hormone (100% Mus musculus) protein inhibitor of activated STAT 3 adult mustilior of activated STAT 3 for inhibitor of activated of the construction of the	RIKEN cDNA 4933433D23 gene 114789 hypohletical protein DKFZp588L0724.1 (75% human) HSPC 166 protein [Homo sapiens] 89 % Interleukin II8
Locus Gene 21807 Tgb 14 22589 Xrp 73205	269378 Ahcy 12814 Cd11a1 76582 50934 Stc7a8	21341 Taflo 73747	67112 Fgf22 11535 Adm	12829 Col4a4 19654 Rbm6 17318 Mid1 20648 Snta1	12508 Cd53 17155 Mania 13731 Emp2 21407 Tcf15 20194 S100a10	66335 Alpévici 54610 The Ida 1256 Cdk2 15227 Fouria 56388 Tspi0 13393 Dk3	66717 78943 Em 1 56316 Ggcx 114568 Kn2-20	77764 229615 Pias3 78651 26563 Rort 16188 II3ra	67554 68979 46501 18183 II8
Chistre Access 9 NM_00936 9 NM_009356 9 NM_009530 9 AKU14175 9 AKU1223	9 NM_016861 9 NM_025640 9 NM_007729 9 AK010877 9 NM_016872	9 AKUT6524 9 NM_021441 9 AKU14309 9 AKU04090	9 AK008922 9 NM_01483 9 NM_009627 9 AF330257 9 AK005253 9 NM_019477	9 235167 9 NM_011251 9 NM_010797 9 NM_024449 9 NM_009228	9 AKO17598 9 NM_007651 9 NM_006548 9 AKO2130 9 AF063878 9 NM_00912	9 NM_025494 9 NM_018775 9 NM_019786 9 NM_010428 9 NM_010428 9 NM_010055 9 RC002239 9 NM_024281	9 NM_025725 9 NM_023913 10 NM_019802 10 NM_025951 10 AY028608	10 L348/6 10 AK020723 10 NM_018842 10 NM_01844 10 AK007241 10 NM_013845	10 AK003286 10 NM_02633 10 BC004785 10 AK007540 10 NM_031168

Locus Gene Description		2	CASS CASS CHARACTER POPEN GOS	25446 DECEMBER CO13 profesion process series administration passes according activates according to the series of the	56532 Riok3	22671 Zfp179	71718	23890 Gor34	7 1897	67863 Sk25a11	27878	17701 Mey 1			76933 TLH29 protein precursor (Homo spatiens) 63 %	Evx1	29	12986 Cs/3r colony stimulating factor 3 receptor granulocyte cs/3r			Cd63	67328 niken cdna 1700038f02 1700038f02rik					68936 ref:NP_478062.1 - chromosome 21 open reading frame 51 (Homo sepiens) 82 %	68147 Nota 1 nudeolar protein famity A, member 1 (H/ACA small nudeolar RNPs)		**	2055 CAPATE CUIDAITE LESS ENOT France servicin & ALCE & Albert chance innecessialism featur & ALOGRE A17 MEEA1 11 M.	Fzd8	Syngri	•	20941 Svp2 A40059 seminal vesicle secretory protein IV precursor - mouse (fragment) 100 %		٠	16328 Immp intranucear protein	SC MD Cross brain other draws mark Ad 28 mm Lesendridad emitted ministra interest interest product		50785 Hs6s11 heparan sulfate 6-0-sulfotransferase 1	Rab6	 Scyct		Kirapa	75040 KKIKEN CONTAK 4530504 NUO BENE		
Cluster Access	10 NM_025788	10 NM 025771	10 A.PBR061	10 NM 030690	10 MM 019955	10 NM 009548	10 AKO04582	10 NM 011823	10 AK009282	10 AK009487	10 BC002307	10 MM 010836	10 NM 016704	10 NM 009124	10 AK010014	10 NM 007966	10 NM_008903	10 NM 007782	10 NM 023197	10 AK005962	10 NM 007653	10 NM 026092	10 AK00/32/	10 AKOOSE 10	10 BC005656	10 NM 025975		10 NM_026578	10 NM 031173	10 AF3/3046		NM 008058	NM_009303	S		10 NM_012065		10 NM 008383	•			 NM_024287	 0	5/6	-,	11 AK015697 7:	,	

Charter Access	5000	8	Description
11 NM 023162	66138		RIKEN cDNA 1110014N07 gene, nuclear RNA polymerase I smail specific subunit [Mus muscatus] 100 %
11 AK0156/3			
10/620 MM 11	, 2002		
EUCESO MIN II			NINEN CLINA SOSUARSILA BENE
11 AF 188286			
NIN CIOLOR			
11 NM 015759			
11 AK006268			
11 BC003249	106628 Trip10	50	thyroid hormone receptor interactor 10
11 L13204	15223 Foxi1	oxj.	hrif-3/forkhead homolog-4 hfh-4 bp 256.,303 activation domain region il 508.,810 winged hefix dna binding
11 KM_007391			
11 NM 021480			
11 AK011413	17840 Mup1	lout	major urhary protein 1
11 AK014671	74503		aniton identified by memorinal antibody MRC OX-2 recentor (41% Mrs muscobs)
	11876 Adn	Ę	riveria
C38000 PNN 11	10 P 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	900	112111111111111111111111111111111111111
DOSON NA I		8 :	una so pou
DOVOTO MIN IT	16835 Ldf	Ħ	low density upoprotein receptor latt
11 NM 024264	74768		A39740 sterol 27-monooxygenase (EC 1.14.14) cytochrome P450 27, precursor - (74% human)
11 AK018638			
11 NM_009427			
11 NM_011219	19283 Ptprz	24	dsd-1-proteogiycan phosphacan homolog
11 NM 019432	-		
11 AKD16475			
BIOLIO MM 11	18412 Saetm1	E E	carriativemo 1
0.5000 100	2226 2626		ייי ביי
SCEND WN II	17 (177	à	rep-02
11 NM_030691	80719 lgsf6	<b>2</b> 10	immunoglobulin superlamily member igsf8
11 AF326960	21380 Tbx1	×	1-box transcription factor (bx1
11 NM 011396	20520 Skc22a5	2205	organic cation/carnitine transporter octn2
11 NM O08353	16161 B12m1	241	injectation (2 persents 15cm)
OCESTO PAN 44		į	
61 010111160			
00011000 111			
II MM OI SEB	1		
11 AF285091	68659		AD021 protein (84% fruman)
11 AK017508			
11 NM_018827			
11 AK006984	76615		S13035 espartate transaminase (EC 2.8.1.1) - human 40 %
11 MM 009405	21953 Inni2	Ę	fast fiber transmin i
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ARUDAZU6	797/9		ALIO TUMAN PICKERI ALCIO (PICKERI COPI IO) (ACCOS) BU PA
11 KM 013463	1960	_	galaciosidase, apria
11 NM 021399			
11 AK004401	71797		chondrolin 4-sulfotransferase (Mus musculus) 47 %
11 Y11717	15400 Hoxa3	ra3	hoxa3
11 NW OTHER	27028 Frman	2	andhinibat mambiana associated protein
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11 8(003/33	ZIUI48 SKSUSE	Mae	Sould Camer rams to Lanc variable to
11 AK016111	75264		RIKEN CLWA 4930553713 pene
11 NM_018819			
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11 AK002581			
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11 NW 022893			
44 MM 007750	13000		ach at months white on white shows \$10011024
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Gene Description	rth4 retinoid x receptor interacting protein rip14-1no.6 alpha fsoform omes tbr2 1-box-containing	ggk N-scen/glucosamine kinase rp8 receptor-activated calcium channel trp? drosophila transieni receptor potemial protein mb3 laminin hela 3.		_		pp1 sphingosine-1-phosphata phosphatase 1; sphingosine-1-phosphate phosphatase [Mus musculus] 100 %	2		progesterone membrane binding protein (73% Homo saplens)			If fro, fringe-tike I (Drosophila)	kinesin kesaw chalin membar kita		DVS27-related protein [Homo sapiens] 51 %	ration (estitin retr	hold chearbedings 25 mount		neslin			a i solule carrier lamily sodium/hydrogen exchanger member tidda 1; done image:3500839	1.1 and the male Reservition of the advanced 2000/12/03	_		CGI-99 protein IHomo scaleral 97 %		RIKEN CDNA 4930542ND7 gene		oold shock domain protein A male germ celt-associated kinase		ref.NP_116187.1 • hypothetical protein FLJ14503 [Homo sapiens] 71.76 %
Locus Gar	20186 NrTh4 13813 Eome	56174 Nagk 26946 Tmp8 16780 Lamb3		73246 Rassfe 20707 Spill	12018 Baki	81535 Sgpp1	18131 Notch3	13067 Cycl	70804		12300 Cacng2	76157 Frd1	16571 Kila	67285 Sdccag10	77125	57284 Retn	DESCE DISCOLU	1301 0000	18008 Nes	12764 Cmas	16511 Kcnh2	20544 Sic8a	71770 An7h1	93687 Csnk 1a1	66230 Mrps28	68045	11982 Atp10a	75185		56449 Csda 17152 Mak		78283
Cluster Access 11 NM_021394 11 AK010015	11 NM_009108 11 AB031037 11 AK020683		11 AK004821	11 AK005472	11 AJ005350 11 NM_007523 11 NM_009047	11 NM_030750 11 NM_030750	11 NM 008716		11 AK014543 11 NM 013822	11 AK010717 11 NM 020514	11 NM_007583	11 AK018094 12 AK016791	12 NM 024179	12 AK014025	12 BC003847 12 NM 011543	12 NM_022984	12 AK012851	12 NM_021600	12 NM_016701 12 RC017625	12 AJ006215	12 NM_013569	12 NM_016981 12 AF302138	12 AKOO731	12 AK019176	12 AK011038	12 AK008080 12 NM 026528	12 AF156549	12 NM 029199	12 NM 009367	12 NM_019638 12 NM_008547	12 AK008928	12 AK019929

Description	ogy menorate skin inter total branch. Lamb endetheldi menter 7 energrad.	RIKEN cDMA 4930560E09 (Mus masculus 25 %)	RIKEN cDNA 1700084J12 gene	Z208 HUMAN Zinc Enger protein 208 53 %			hypothetital pratein FL/20764 (73% human)	Mus muscutus 0 day neonate skin cDNA, RIKEN full-langth emiched library, done:4632413E21 producti-weakly shrikar to PMOSPHOLIPASE B I Pathus norveptous)	retinal degeneration 82 homidog (Orosophila)	PET112-like (yessi)	guanyate nucleotide binding protein 2	kv3.3 potassium chammel protein exon	peraxisomal biogenesis factor 3	GLE1 RNA export mediator-like (yeast)		very tow density tipoprotein receptor
Locus Gene	72324 Tem7	72747	73486	170763			71963	78281	19679 Rdgb2	229487 Peri 121	14469 Gbp2	16504 Kenc3	56535 Pex3	74412 Gle11		22359 VIds
Chaster Access	12 AKO10361	12 AK013267	12 AK006993	12 AF242377	12 NM_026047	12 AK009010	12 8C012953	12 AK019494	12 NM_011256	12 BC005709	12 NM_010260	12 NM_008422	12 NM_019961	12 AK016671	12 M36654	12 NM 013703



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6.3988 Gmfg gila maturation factor, gamma
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752 74209 Rabbe-francing potent 2 (in procedure)
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hydelit CL28B protein -rat (31 % R.novegous)
hypothetical protein MCC15435 (Homo sapiens) 51 %
CD9 antigen
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72162 Ddx36
14089 Fap
14836 Gsc
56447 Copz1
70729 Capon
69922 Vrk2
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Mus musculus 10 days embryo whole body cDNA, RIKEN full-length entithed library, chore;2910034818 producthypothelical protein RAP2A, member of RAS oncogene family (K-rev); K-REV [Homo sapl 39 % target of myb 1 homolog (dribchen)
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durbs NADH dehydrogenese (ubquirone) 1 beta subcomplex
durbs NADH dehydrogenese (ubquirone) 1 beta subcomplex
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hypothetical protein ELLZ3467 [Homo sapters) 39
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Huringtin intereding protein K; hypothetical protein [Homo sapters) 39
Huringtin intereding protein K; hypothetical protein [Homo sapters) 39
Huringtin intereding broadh protein Clot 100 precursor (89% human)
LGAQD HUMAN LG SarbAd Associated Sm-like protein LSm3 (MDS017) 100 %
HREN CDNA 620563041 1 gene
HREN CDNA 620563041 1 ge
                                                similar to pic. 100322 - 100322 hypothetical protein KIAA0542 - human 62 %
BR21.2 Liste 12 (profine irch); Bci-2 like profine-rich protein 12 [Homo saplens] 81 %
unc-34 homolog A (C elegan)
sp.093189 - KHL2. HUMAN Keich-like protein 2 (Actin-binding protein Mayven) 89 %
nuclear factor of kappo light polypeptide gene enhancer in B-cells inhibitor-like 1
mannosidase 2, alpha 82
mannosidase 2, alpha 87
mannosidase 2, alpha 87
mannosidase 1, alpha 87
mannosidase 1, alpha 87
mannosidase 1, alpha 87
mannosidase 2, alpha 88
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mannosidase
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anityrin repeal and SOCS box-containing protein 12
dishevelled 2 dsh homdog drosophila 0vi2
cysteine and historie-rich domain (CHORD)-containing, zinc-binding protein 1
RuvB-like protein 1
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Sestrin 3 (100% Mus musculus)
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solute carrier family 35 member stc35a2 udp galactosa translocator; mugt1 udp-galactose transporter
Similar to Protein transport protein Sec24D (SEC24-resited protein D/g32% Human)
polycythemia rubra even 1; cell surface receptor; cell surface surface; cellon s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transcription elongation factor B (Sitt), polypeptide 3
polymearse (RNA) II (DNA directed) polypeptide G
prosphodiesterase I A. calmodulin-dependent
RNEN cDNA 1790025602 gene
RNEN cDNA 1700025602 gene
RNEN cDNA 1700025602 gene
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inhibitor of kapoe light polypeptide enhancer in 8-celfs, kinase complex-ossociated protein
suitatises 2
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Similar to WD domain, G-beta repeat-containing protein [Homo saplens] 63 %
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peroxisone bogenesis factor 1
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hysy oxidasa-related profein 2 loc2 loc2
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RIKEN CDNA 2010001E11 gene
pleckstrin and Sec7 domein protein [56% Homo sapiens]
RIKEN CDNA 2310079NX2 gene
13 days embryo head riken cdna chone:3110031013
ras homotog gene family, member H
low density lipoprotein recoptor-related protein 1
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16971 Lrp1
19730 Ralgds
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COURSE ACCESSES

B NIM_2056239

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Na macuta suppression of tumoripentally suppression of tumoripental suppression of supp
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acidic epididyrral pycoprotein 2
vascular Rab-ASPTREC-constituting BUBZ-like protein 1 [Mus musculus] 48.54 %
vascular Rab-ASPTREC-constituting BUBZ-like protein 1 [Mus musculus]
vascular Rab-ASPTREC-constituting bubz-like protein 1 [Mus musculus]
rucin 10, submandibular gland salivary mucin
hypothetical protein FL1,1326(17 % human)
C-type (coliculus dependent, carbohydrate-recognition domain) lectin, superfamily member 5
microtubule-associated protein 1 light chain 3 alpha; MAP1 light chain 3-like protein 1; microtubule-associated proteins 1 A/18 light chain 3 | 100% Homo saplens].
RIKER CDN4 170012/203 gratum-dependent, delia Isoform i
givalent calion tolerant protein CUTA [Homo saplens] 91 %
        Describtion
Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:4921510.117 product.hypothelical EF-hand containing protein
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                                                                                                               beaded flament structural protein in lens-CP94
RikEN cDNA 2310043108 gene
eukaryotic translation inilialion factor 3, subunil 4 (defta)
cycin-dependent kinase 5
Array repair complementing defective repair in Chinese hamster cells 2
Son cell protiferation protein
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DNA segment (Dr. 18 ERATO Dat 465, expressed
NNA segment (Dr. 18 ERATO Dat 465, expressed
Nypotheriteal protein FL2012S (83% Homo saptera)
HSPC134 protein (60% Homo saptera)
RIKEN cDNA 5630411,007 gene
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RIKEN cDNA 4933440J22 gene
chemokine (C-X-C motif) ligand 5
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11 NM_ 001531

11 NM_ 0016379

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11 NM_ 001633

11 NM_ 0016324

11 NM_ 0016364

12 AKOD5884

12 NM_ 001991

12 AKOD589

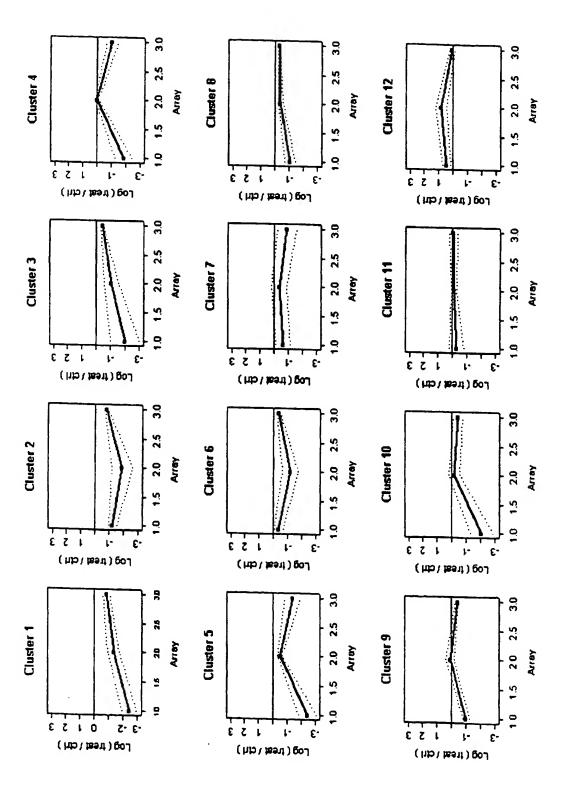
12 NM_ 001991

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67711		RIKEN CONA 2510027N19 [Mus musculus]
78283		rei:NP_118187.1 - hypothetical proteth FLJ14503 (Homo sapiens) 71.78 %
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13618	Edut	endohlein-b receptor edurb
13813	Eomes	(br.2 )-box-conflairing
67111		Similar to PLT_HUMAN Protein PLT 81%
18103	Nme2	expressed in non-metastatic cells 2, protein
17918	Myo5a	myosin Va
68991		HSPC182 protein [89% Homo sapiens]
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7573		RIKEN CHAN 14 20700M05 gene
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14538		glucosaminyfitansferase, i -branching enzyme
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1		
	20348 67711 78283 22310 13818 13813 67111 68981 17918 67121 12353 14539 14539 17283 16538 17283	83 V2r4 111 813 Ednrb 113 Eomes 114 Ednrb 115 Eomes 117 Eomes 118 Ednrb 119 Eomes 119 Environ 119 Eomes 119 Environ 119 Eomes 119 Environ 119 Environ 119 Environ 119 Ednrb 119 Environ 119 Enviro

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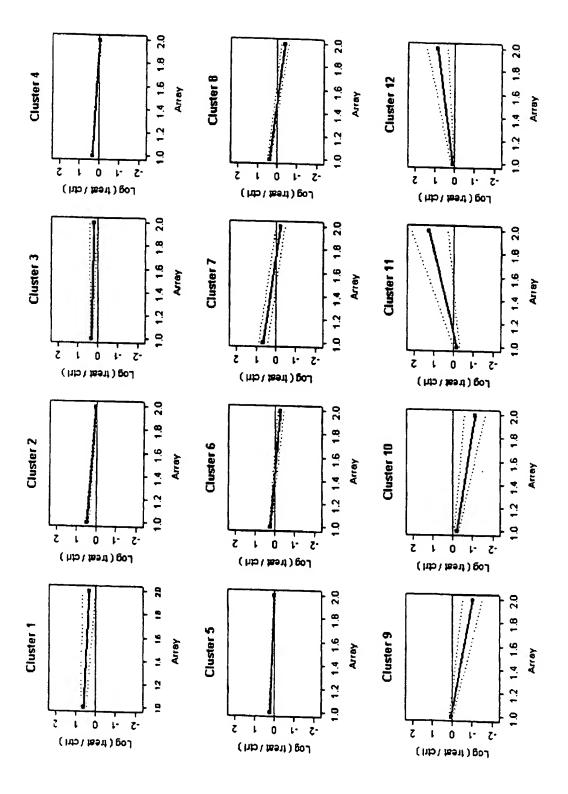
10 NM_01189

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11 AKOO2931
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11 NM_052993	70207		
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11 NM 013592	17183	Matre	marine and the second
11 NM 008217	15116	Has3	hyeluroran synthese 3 has 3
11 NM 025658	20999	MsAbdd	membrane-spanning 4-domains, subfamily A, member 4D
11 NM 008297	15500	Hsf2	heal shock factor his?
11 NM 02/B10			
11 AKUUSSE	73533		RINE NOTATION OF THE PROPERTY
11 AKO10480	29/9/		HINEN GUND ZA 100 131Z gene
11 AKUTSUDI	56733	9	Small for polassium voltage-gated channel, subhamely Co, member 1; polassium channel KH2 (78%, Homo saptens)
SCHOOL ON THE	836/3	Ē	camerious care
11 BC003810	28507	reg.	emplyonic estimated give
11 NM 018280	64613	Alpevicz	A I vasa, He introducing, V 1 student L. Bottom 2
11 MM 016780	21000	cdire	Secrete and Cazio-Create Sequence Protein 5
10 MA_009757	55127	c de	grown outervisions actor-be gati-sp. cone morphogenetic protein 15 pmp15 outers actor-be gati-sp.
11 64012248	0//6/	200	NINCH CAVA 465.50 ZANI 5 GOTE
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11 NM 023784	75581		SINCE ADM 2010 File II the researches 100 %
11 NM 009343	21652	Ē	PHD (inner protein 1
11 AK012908	212127		RIVEN COMA 2810008.04 perse
11 AK014404	74014		RIVEN CONA 3732407.023 gene
11 NN 013887	30044	Oprid	ossin 4 (melanousin)
11 NM 021421	52477		DNA segment, Chr. I. ERATO Doi 399. expressed - RIKEN cDNA 2610307121: hypothetical protein, MNCD-4273 liMus musculus! 100 %
11 NM 013714	26573	Ireb(1	interferon response element binding factor 1
11 AKG07777			
11 AF318301	192198		CGI-74 protein; CGI-69 protein (Hamo sapiens) 88 %
11 AK019409	103425		RIKEN 6JNA 3100002P13 gene
11 X65508	16873	Kr1.5	type i kersitn mbr a-1
11 NM 007875	13478	Opagt2	dolichyl-phosphale apta-n-acetylglucosammephospholnarsfense 2 dpegi2
11 NM 021454	58801	Pmelp1	photoel 2-mytalsel-3-acetate-induced protein f; Nova protein [Mus muscutus] 100 %
11 AK02/182	7/616		KINEN CASSONERS IN SECTION OF THE CASSONERS OF THE CASSON
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11 AK007269	0350		ום תפופ מוווים אל מוווים מווים מ
11 NM 007892	13558	E26	e2) travertation factor 5 done mac:6043; e21:5 tratein
11 NM 010298	14658	e e	aly dire receptor, beta such unit
11 NM 008868		Lypla1	Ayoutospholipase 1
11 NM_008287		Hrsp12	heat-responsive protein 12
11 NM_009252			
11 AK020464	77323		similar to pp.P11359 - POL2_MOUSE Refrondus-relation POL polyprotein (Contains: Reverse increatainse) Entonucleasej 43,28 %
11 NM_009792			
100070_WN 11			
	77757	200	protein (virginis protein) (1951.) MANT Of I Mooth between their first telebrid medic embrorate (77% M miscalis)
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Chatter Accesses		Pigdr	Ogine Description Pigot prostagandin D receptor
12 NM_009843	12477	Cita	cytotoxic T-tymphocyte-associated protein 4
12 NM_010946	18203	CBZ	P-terminal asparagine amidohydrobase rtan t
12 NM 020515	57272	Ora 16	gene for odorant receptor A16
12 026047	18700	e gid	phosphalidylinosid gyran, class A
12 AND 11365	9890	į	KIKEN GJAZ ZAJOŽIVOJ POPIN
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12 AKD14783	7306.4	57.5	Must material south, Kilkert Indestigning and Indestigning Come; 1/00/22/CU7 product sementheenthe kingse 33,
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12 MM 000433	32004	בר ב	open recording to the control of the
12 NW 008536	62033	Todat	Injugación de la contractor de la contra
12 AKO14791	7	- Safe	Haismemorane & superiarmy member 1
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12 NM 011328	20.00	3	or Ori (amazeni amarina) ino a
12 AK015166	74907	š	Socient intermedials flammed binding periods 6,0000 lidoses seedand 25, 24 ft.
12 AK015780	74717		Process research seasons and process, John Spring S
12 NM 007900	13605	Fet2	POLICE TO THE POLICE TO THE POLICE THE POLIC
12 NM 020605	57340	1	over controlled to the mendered 100 %.
12 AK004401	71797	}	condition 4-subcitaristense (As meeting 47%
12 AK016572			
12 AK012399			
12 AF333770	170571	Critrap4	contact in associated protein 4
12 MM_011888	26888	Clecs f8	C-type (calctum dependent, carbotydrate recognillon domain) lectin, superfamaly member 8
12 AB036341			Mus musculus mRNA for CN 8 scFv, complete cds
12 NM_010498	15931	\$P\$	kduronale sullalase ids
12 AK019494	78261		Mus musculus 0 day neonale skin cDNA, RIKEN full-length enriched library, chne:4632413E21 product weakly similar to PMOSPHOLIPASE B [Rattus norvegicus]
12 NM_009796	12339	Capn7	calpan 7
12 AK006511			
12 AF264049			
12 NM_021453	58803	Pep	18 days embryo riken cdna done: 1100036a17; pepsinnegen pepi aspanito proteinase
12 NM 023045	65246	Randp 16	ran binding protein 18
12 BC018550			
12 AK006187			
12 M65237			
12 AK01/277	;		
12 BC004678	101333		expressed sequence AW123240
12 AK010429	71956	1	Z147_MOUSE Zinc (Inger protein 147 (Estragen responsive Inger protein) (Elp) 32 %
12 AK019498	15507	Hsp25	day neorate skin riken odna deone: 462,415,09
12 NM_0084/9	16768	Leg3	∳mphocyfa-ectivation gene 3
12 MM 03536			
12 AKON7826	68530		DIKEN ATMA 1110702 2 2 AMANA
12 MM 000396	21928	Tridain	times recovered index of his state of the st
12 AK005661			
12 AK004688			
12 NM 009799	12346	Ş	carbonic anhydrase i
12 AK017508			
12 NM_008298	15502		DnaJ (HspdO) homolog, subfamily A, member 1
12 AJ278787	110880	Scride	
12 AK014671	74603		antigen identified by monocional antibody MRC DX-2 receptor (41% Mus musculus)
12 NM_016884	15381	Hrepc	helengelenbus nudder fiborucieopinden o harripo
12 AK011843	70439		TAFTS RNA polymerase II, TATA box binding protein (TBF)-essociated factor
12 AK014813	42835	,	SVT3 MOLISE Consulcidentation II (SAII) 38 %
12 NM 027719	71202		FIREM 2014 49343EE Of Mar meeting 100 %
12 AK013889		Wdr17	WD repeal domain 17
12 AJ278890	192120		B-box and SPRY domain contributing

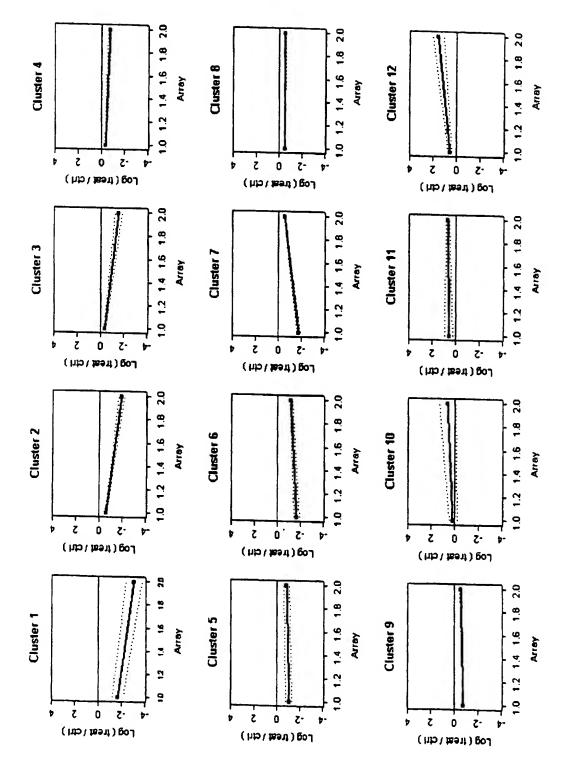
Children Access Copies Description 12 BOOT 12	RRIEN CONA 48124 1272 gene	KINEN CATA AS INVASION FINEN EMBACKATORY OF THE STATE OF	hypothetidal protein MGC11257 (82% human) KIAA 1074 protein Huma samiena (45.33 %.	ubiquin specific professe 20	tuberin-like protein 1	RIKEN cDNA 4933433023 gene	JC4 131 gloma pathogenesis -ekiled protein - human 40 %		ltyrotrophin-releasing hormone receptor trh-t: thyrotropin releasing the		ephin AS
- Oems				Usp20	Tulip				Trh		Elna5
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Cluster Access	12 AK014588	12 AK002512	12 AK014271 12 AK015473	12 AK006800	12 AK009111	12 NM -028232	12 AKDO7167 12 AKD07167	12 AF220134	12 NM_013696 12 AK016419	12 NM 025459	12 NM_010109



	Description Calson Lestin 2		_	major vault protein	RIKEN cDNA 8430416G17 gene	hepatocyte growth factor	regucalcin	Akr1c6		RIKEN cDNA 4930415F15 gene	amyloid beta (A4) precursor protein-binding, family A, member 3		inhibitor of kappaB kinase gamma	•	Eaf1 protein	WNT1 inducible signaling pathway protein 1	nuclear RNA polymerase I small specific subunit [Mus musculus] 83 %	:yypothet,ical protein FLJ10694; hypothetical protein FLJ10889 [87% Homo sapiens]	prolactin-like protein E	hypothetical protein, 12H19.01.T7	transcriptional adaptor 3 (NGG1 homolog, yeast)-like	histocompatibility 47	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsiton		enhancer of zeste homolog 2 (Drosophila)	regulator of g-protein signaling 9-2 isoform rgs9 rgs9-2 isoform striatal-enriched alternatively spliced product	syntaxin binding protein 4	ribosomal protein S6 kinase	trehalase (brush-border membrane glycoprotein)	breast cancer metastasis-suppressor 1	keratin associated protein 13	Similar to mammalian inositol hexakisphosphate kinase 2 [Homo saplens] 91 %	olfactory receptor 67	expressed sequence Al448222	EH-domain containing 3	Similar to amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 2 [50% Human]	POU domain, class 3, transcription factor 2	prolactin like protein I	ubiquitin carboxy-terminal hydrolase L1	pellino 1- ref:NP_075813.1 - pellino 1; RIKEN cDNA 2810468L03 gene [Mus musculus] 100 %	myosin IB
-	Gene Caso?	Puma-g	ဗွ	₩		Ξĝ	Rgu	Akr1c6	Tpx1		Appa3	mor17-1	Ikbkg	<b>Olfr159</b>	Eaf1	Wisp1			Pripe		tada31	<b>14</b>	Nfkbie		Ezh2	Rgs9	Stxbp4	Rps6ka4	Treh	Brms1	Krtap13		Olfr67		Ehd3		Pou3f2		Ξ	_	Myo1b
***************************************	Logus 12373	80885	14836	78388	71469	15234	19733	83702	22024	73862	57267	209102	16151	29849	74427	22402	76416	101437	19113	80982	101206	109815	18037		14056	19739	20913	56613	58866	107392	16699	76500	18368	52504	57440	72149	18992	67000	22223	67245	17912
	Gluster	1 NM_030701	1 NM_010351	1 NM_080638	1 AK018420	1 X72307	2 NM_009060	2 NM_030611	2 NM_009420	2 AK015144	2 NM_018758	2 AF133300	3 NM_010547	3 NM_019476	3 AK016628	3 NM_018865	3 AK006236	3 AK014260	3 NM_008930	3 NM_030728	4 AK003405	4 BC011091		4 AK020739	4 NM_007971	4 NM_011268	5 NM_011505	5 NM_019924	5 NM_021481	5 AF233580	6 NM_010671	6 AK005166	6 NM_013619	7 AK018444	7 NM_020578	7 AK011451	8 NM_008899	8 NM_025896	8 NM_011670	9 AK020915	9 NM_010863

Gene Description Cab140 calcium binding protein	inorganic pyrophosphatase [Homo sapiens] 71 %	odorant receptor S1 gene (Mus musculus) 100 %	gap junction membrane channel protein alpha 10	FXYD domain-containing ion fransport regulator 7	methyl-CpG binding domain protein 3-like		elk4 member of ets oncogene family clone image: 3589378	AFG3(ATPase family gene 3)-like 1 (veast	immunoglobulin superfamily containing leucine-rich repeat	G protein-coupled receptor 90	claudin 2	olfactory receptor 64	FYVE and coiled-coil domain containing 1	S12207 hypothetical protein (B2 element) - mouse (68% human)	cystatin 8 (cystatin-related epididymal spermatogenic)	G-protein-coupled receptor 50
5000		Ors1	Gja10	fxyd7	Mbd3l	Zfp354c	EIK4	Afg311	ıs	Gpr90	Cidn2	Offr64	Fyco1		CstB	Gpr50
Logus 12282	74776	56858	14610	57780	73503	30944	13714	114896	26968	80978	12738	18365	17281	71546	13012	14765
Charles Access 9 NM 021395	10 BC011417	10 NM_020288	10 NM_010289	10 NM_022007	11 AK018931	11 NM_013922	11 NM_007923	11 AK012394	11 NM_012043	11 NM_030726	11 NM 016675	12 NM_013616	12 AK013060	12 AK018561	12 AF090691	12 NM_010340





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Description  RIKEN CDNA 0610020102 gene  syntrophin, addit 1 sec61 alpha subunit 2 s. cerevisiae done mgc:6359; sec61a2 storndtin-like protein 1(79% human)	proprotein convertase subtilisin/kexin type 4	ENC1, MOUSE Ectoderm-neural cortex-1 protein (ENC-1) 80 % / RIKEN CDNA 2700023J09 gene RIKEN CDNA 120001118 gene	politivity receptor feater 5 - the feature of the	Janus Miase 2	CAMP (Mus musculus) 100 %	Ras association (RaiGDS/AF-6) domain family 6 S508S3 translation releasing factor eRF-1 (99% human)	NK1 Iranscription factor related, locus 2 (Drosophila)	chronosome 20 open reading frame 108 [Homo sapiens] 77 % cut drosophila-like 2 cut2 cut drosophila-like 2 cut2 T 125 15 hypothetical protein DKFZp4348103.1 • (26% fruman)	chloride intracellular channel 5	RIKEN cDNA 2310002F 18 gene Similar to MOUSE Alpha-actinin 3 (Alpha actinin sketetal musde Isoform 3) (F-actin cross linking protein) 28%	carboypeplidase d cpd	centromere autoantigen C solute comier family 7 (cationic amino acid transporter, y+ system), member 10	serineIhreonine protein kinase SSTK RIKEN cDNA 2310034L04 gene	Mus musculus 18-day embryo whole body cDNA, RIKEN full-length emiched library, clone:1190009E12 product:hypothelical SOCS domain, C-terminus of STAT-Inhibitors containing profein florobiast growth factor 1	Mus musculus Tni receptor-associated factor 3 (Trai3) gene, partial sequence; and armioniess precursor protein (Arm) gene 130249 celi protiferation antigen Ki-67 - mouse 100 %	C Iq refaled factor catsyntenin-1 protein		leukotriene od synthasse RINCIN CDNA 4933425k0 gane	supplicative dec 2.5 s. Joan iv 1954 mus musculus) 2.5 oligodeemises 16 Text ( RR F I)-Interacting nuclear factor 2
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9 NM 024264	74768		A39740 sterol 27-monooxygenase (EC 1.14.14) cytochrome P450 27, precursor • (74% human)
9 AK005168			
9 AK004171	68792		sushi-repeat protein [Homo sapiens] 93 %
9 NM_018804	54527	Syli	Synaptotegrafin xi
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9 AK013274	19087	Prkar2a	protein kinase, cAMP dependent regulatory, type it alpha
9 AK016052			

e Description	a20 solute carrier family 25 (mitochondrial carnillne/acytcamiline translocase), member 20		r III macrophage inflammatory protein- f alpha receptor-like mip-1 receptor like-1 transcriptional co-activator with PDZ-binding motif (TAZ) (91% Homo septens)			Ans muscutus adult male tests CDAA, RIKEN tul-tentiched library, clone:4930448A20 product undassifiable				25 COT'N (CONTINUING PROTOTOR) PROTOTOR, BUDURII 9 Antherio Ast		_	NAP kinasa-edivided in kinase 2  - Manager advantant kinase 2		_	lissue inhibitor of metaltoproteinase 1	phistore h2a.z		banscription factor like 4 tota	d mitogen-activated protein kinase		MUST LIBOURDS SOURCE TO THE TOTAL OF THE TOT		eduli mele legis ikter odne domes/95437105 eduli mele legis ikter odne domes-asa R Innestriation factor 3) (AB 63% Max musculus) Transitation factor BTC4 (DNA ochamesas R Innestriation factor 3) (AB 63% Max musculus)	- 5	-		2 zinc imger protein 162	oliacioy receptor 6		CONF INC. 200 CONF. CONF			_	_	Ī		HAREN CUMA AUGUSANDO GENE At mercalise DRF1 and ORF2 nemed	Must musculus adult male testis CDNA, RIKEN Mill-length enriched library, done:1700041N15 product: CHEMOKINE-LIKE FACTOR 2 VARIANT 2	hypathetical protein FLJ12783 [Homo saplens] 96 %	polymerase (DNA directed), sipha i	mita program wood vegoda protein coloran price foreign 3 (mean ideoda)	Coory Sundament Section 5 (granteer) (1908)	
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## FIGURE 10-11

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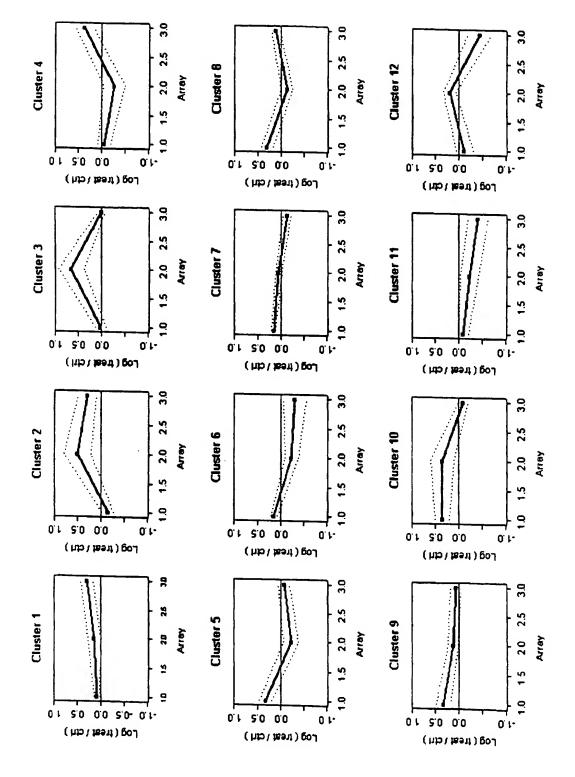
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## FIGURE 10.12

Chatter Access	Locus	8	Description
	56363	Trneff2	transmembrane protein with EGF-like and two follistations of proteins 2
12 NM 023397	67881	1	magnesum-dependent phosphalase-1   mus muscuus) 100 % Kinch CONA 1010004RZU gene
12 AK007552	56036	25 5 5	ordin I.
12 AK011545			
12 AK010648	70024		Similar to MCM to minichromosome maintenance deficient 10 (5. cerevistae) [81% Homo sapiens]
12 BCUD4801	15220	Forot	fort head transcription factor hth-11 member of but-3/fork family, hth i
12 AKO11615	71517		esophageal cancer associated protein [Homo sapiens] 92 %
12 AK014861	70894		Mus musculus aduli male testis cDNA, RIKEN full-length enriched library, cone:4921510J17 producthypothetical Er-hand containing protein
12 NM 025725	2119	i	RIKEN CONA 42215 13E08 (Mars muscutus) 100 %
12 NM 012032	26943	Tde 1	merubane professional units and the southern to be a demander of the southern release of the following the southern them seems of the southern the s
12 NM 023210	66471	S	manp-lanp-like protein recens not account. To it days enter you men consistency consistency and account of the consistency of t
12 NM 029231	75273		RIKEN cDNA 4930563C04 gene
12 AK021021			
12 AK009634			
12 NM_021524			
12 NM_0244B0			
12 AK002842			
12 AK013518		4	firmblas annuals forder & field & drie arren
12 NM_010203	141/6	2	includes by the control of the contr
12 NM 023224	8 20 20		ability the line in the contract of the contra
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12 AK019929	78283	51-745	FBL.RT. 10 10.1. Inflyenmental production of the second production of the second secon
12 NM 009521	52415	SHA S	Willy season min to integrate on the control of the
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12 NM_008493	10040	9	inda
12 BC004151			
12 NM C1	26549	Red (bo2	injegrin beta 1 binding protein 2
12 NM 015797	50762	Fbxo6b	F-box only protein 6b
12 NM 008052	14357	Š	delicx1
12 NM 019485	56014	Olfr70	oliaciory receptor 70
12 NM 008838	18701	P	phosphatidylinositol glycan, class F
12 NM 010145	13849	Ephx1	epoxide hydrolase 1, microsomal
12 BC008101	68939		hypothetical protein MGC2827 [94% Homo sapiens]
12 NM 026252	67579		RIKEN cDNA 4930447D24 (Mus musculus)
12 NM_010740	17064	Clart	complement component 1, q subcomponent, receptor 1
12 NM 027884	21961	Tris	lensin
12 NM 025443	66249		RIKEN CDNA 1810003NZ4 gene
12 NM_016737	20867	Slip1	stress-induced phosphoprotein 1
12 NM 008821	18630	Pelz	pleanetyfora expression named a second of the second of th
12 NM_008968	19223	SB	prostagament iz prostacymin synunciae prijas
12 NM 025331	0,100	6.100	the state is explore voiced with a he insertion containing stop codon.
12 050366	19970	Sam	August Reports as a specific vision in the second s
12 ANUMADO	73030	Mercan	Alderonalist measurement (No. 1974)
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12 NM 023033	76647	1	RIKEN CON 170002692 cene
12 ANUG362	19184	Period	professe (prosome, macropain) 26S subunit, ATPase 5
12 NM 016289	53324	Notx2	neuronal pentraxin 2 nptx2: np2
12 NM 031255	83434	Rsh11	radial spokehead4 protein rshill
12 NM 013799	11907	Alet	arginine-Ima-protein transferase ste 1
12 NM 026639			
12 AK018698			
12 NM_018798		1	e de la companya de l
12 [38281	16365	6	יוו מיניים באריים איניים באריים באריי
12 NM 080557	69150	Snx4	sorting nextn 4 (Aus musculus) 100 %



NM 080450

Charter

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professome (prosome, macropain) subunit, eighe type 3
A28438 lactoferrin precursor - mouse
Mus musculus adult male testis CDNA, RIKEN full-length enriched library, clone:4921504102 product:hypothetical Serine-rich region containing protein
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Mus musculus 0 day neonate head cDNA, RIKEN full-length enriched library, ctore:4833422M21 product:hypothetical protein
17 days embryo haad riken cdna ctona:3222401p09; atp-binding cassette protein abcb8
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guanine nucleotide binding protein alpha transducing 2 gnat2; adult male tongue riken cona clone.2310018g23
Son celt profiteration protein
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pellino 1-ref.NP_075813.1 - pellino 1; RIKEN CONA 2810488L03 gene [Mus musculus] 100 %
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10 days embyo riken cdna clone:28 10003106; nma novel transmembrane protein
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olfactory receptor 159
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66765 Similar to myeloidflymphold or mixed-lineage leukemia 2 (41% Human)
258811 MOR1714 difactory receptor MOR1714.6
105017 T46427 hypothetical protein DKFZp434G0825.1 (38% human)
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67245 Peli1
20965 Syn2
28401 Map3k1
70235
28913 Zh285
78734 Lypla2
116914 Sic18a2
77145 Wdy3
57442 Kcne3
66402 Sin
19167 Psma3
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14886 Gnat2
20658 Son
20853 Def8
20321 Sdh2
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17775 Nrap
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AK008679
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£ 40-00		5 8 8 m	stimulated by retholic acid 13 Interferon-garman (muith-garman) A k thrase (PKKA) anchor protein 8 A k thrase (PKKA) anchor protein 9 A seleno-gration associated protein PH (mmo sapiens) 88.64 % Contactin associated protein-like 2 A STANDA seleno-gration associated protein PH (mmo sapiens) 88.64 % A seleno-gration associated protein PH (mmo sapiens) 88.64 % A seleno-gration associated protein anchor 1 k (PKKA) A seleno-gration associated protein 1 k (PKKA) A STANDA STANDA STANDA PKECURSOR (FERROXIDASE) (96% Mus muscufus) CERU MOUSE CERU (POLASMIN PRECURSOR (FERROXIDASE) (86% Mus muscufus) B days embryo riken dana done: 1 110032016 B days embryo riken dana done: 1 110032016 COPP (constitutive photomorphogenic) homotog, subumit 3 (Arabidopsis theliana)

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Mus musculas 13 days embryo head cDNA, RIKEN Mil-length enriched library, done:3110003A17 product:unknown EST S100 cark-tum binding protein A8 (catgranulin A) chromodomalin protein. Y chromosomal-film and chromodomalin protein. Y chromosomal-film complex focus (Guanine nucleotide binding protein apha stimulating) complex focus RIKEN GDNA 1110014N07 gene, nuclear RNA polymorase I small specific subunit [Mus musculus] 100 %
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Mus musculus adult male testis cDNA, RIKEN full-length enriched library, done:4933413N12 product.undassiliable
Mus musculus RIKEN cDNA 0610030G03 gene, mRNA (cDNA done MGC:11893 IMAGE:3598444), complete ods
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Tax1 (human T-cell leukemia virus type I) binding protein 1; tax1-binding protein [Homo sapiens] 80 %
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Z.147 MOUSE Zinc finger protein 147 (Estrogen responsive finger protein) (Etp) 32 %
histone cell cycle regulation defective interacting protein 5
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JC6547 high sufur proteh BZE - rat 37 %
LSM3_HUMAN Us srRNA-essociated Sm-like protein LSm3 (MDS017) 100 %
NADH dehyfongenase (ubriguinane) 1 beta subcomplex
gamma-aminobufyric acid (GABA-A) receptor, subunit gamma 1
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                                                                                                                                                                                                                                                                                                                                                                                                                    protein phosphatase 1, regulatory (inhibitor) subunit 3F [83% Homo sapiens] inner membrane protein, mitochondrial RAB3B, member RAS oncogene family
                                                                 Ca 2*dependent activator protein for secretion opsin 1 (cone pigments), medium-wave-sensitive (cotor blindness, deutan) ubiquitin-specific protease ortubain 2 (94%)
                                              profeasome (prosome, macropain) 26S subunit, ATPase, 4
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RIKEN cDNA 2610208C17 gene
keratin associated protein 13
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52440 O6Erd772 |
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14683 Gnas
66136
14489 Mtpn
27204 Syn3
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AK012535
Cluster Access
6 Y12229
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Mus musculus adult male corpus etriatum cDNA, RIKEN full-length enriched library, clone:C030048J01 product:hypothetical protein
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ref.NP_543020.2 - cystatin 11, isoform 2 precursor; cystafin-related epididymal-specific protein; SC13delta [Homo sap 37 %
                                                                                                        twist gene homolog 1 (Drosophia)
multo deduced amino acid sequence of is homologous to c. elegans putative drag protein z73102 b0035.14. homolog
Nit protein 2 (89% Homo sapens)
RIKEN cDNA 20 to 107K23 gene
antigenic determinant of rec-A protein
low density (lipoprotein-related protein 18 (deleted in tumors)
ATP-binding cassette, sub-family A (ABC1), member 1
RIKEN cDNA 57304 17817 gene
GTP binding protein 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           odd Oznan-m homolog 2 (Drosophila)
UDP-CichtAccredica hera-1,3-N-acchtiguosaminytransferase 3
UDP-CichtAccredica hera-1,3-N-acchtiguosaminytransferase 3
UDP-CichtAccredica hera-1,3-N-acchtiguosaminytransferase 3
UDP-CichtAccredica hera-1,3-N-acchtiguosaminytransferase 3
UDP-CichtAcc HUMAN Dynactin complex 50 kDa subunit (50 kDa dynein-associated polypeptide) (Dynamtin) (DCTN- 94 % mannosidase 2, alpha B2
Similar to KHL1, MOUSE Kerchlike protein 1 30% bore marrow stronal cell artigen 1
BCL2-tike 12 (proline rich); BG-2 like proline-rich protein 12 [Homo sapiens] 81 %
                      similar to A49307 98K GTPase-activating protein ABR, brain - human 29 %
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BRF2, subunit of RNA polymerase III transcription initiation factor, BRF1-tite
RIKEN cDNA 9230106L14 gene
spindlin
                                                                                                                                                                                                                                                                                                                                                                                                                                          Similar to hypothetical protein MGC 16309 (Homo saplens) 70 % trehalase (brush-border membrane glycoprotein)
                                                                                                                                                                                                                                                                                                                                                   Similar to hypothetical protein FLJ20548 (82% Homo sapiens) RIKEN CDNA 2610507L03 (Mus musculus) 100 % Similar to hypothetical protein FLJ11259 (Homo sapiens) 93 %
                                                                                                                                                                                                                                                                                                         serine (or cysteine) proteinase inhibitor, clade E, member 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mudix (nucleoside diphosphate linked molety X)-type motif 7 inhibitor of growth family, member 1 pheromone receptor V3R9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DnaJ (Hsp40) homolog, subfamily B, member 7 hypothetical protein FLJ20048 [Homo sapiens] 78 %
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B lymphoid kinase
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                                          RIKEN cDNA 2310016G11 gene
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chemokine (C-C motif) ligand 4
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18368 Olff67
22180 Twist1
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16588 Kin
1303 Abca1
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20720 Serpine2
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74589
12182 Bst1
75736
57257 Vev3
77145 Cut2
57810 Cdon
12143 Bit
16709 Kin1
67528 Nudt7
26356 Ing1
861010 V3R9
57755 Onajb7
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77053 Unc84a
23964 Odz2
72297 B3gnt3
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12828 Col4a3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12944 Crp
14232 Fkbp8
17936 Nab1
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               8 AK004654
8 AK009387
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8 NM_011658
8 NM_023175
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8 NM_05378
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D AK004552
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9 AF343752
9 NM_011856
9 NM_028189
9 AK013799
9 AK013702
9 NM_01850
9 AK013702
10 NM_01852
10 NM_01822
10 NM_018139
10 NM_018139
10 NM_01818
10 NM_01883
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9 AK010876
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Chister Access	Locus	Gera	Description
	70729	Capo	C-terminal PDZ domain ligand of neuronal nitric oxide synthase
	74420		RIKEN CDNA 4933406P04 gene
11 NM_009237	20675	Sox3	SRY-box containing gene 3
11 AK004139	68735		mitochondrial ribosomal protein S18C; CGI-134 protein; mitochondrial ribosomal protein S18-1 [Homo 78 %
11 NM_015733		Casp9	caspase 9
11 NM 021310	57748 Jrmy	Ę	function-mediating and regulatory protein
11 BC006048	14194 Fh1	Ē	fumarate hydratase 1
11 NM 025310	56095	Epcs3	ectoplacental cone, invasive trophobiast glant cells, extraembryonic ectoderm and chorion sequence 3
11 NM 020578	57440	Ends	EM-domain containing 3
11 AF233580	107392	8mms 1	breast cancer metastasis-suppressor 1
11 NM_011622	21968	Tom	target of myb1 homolog (chicken)
11 NM 011783	23795	Agr2	anterior gradient 2 (Xenopus Iaavis)
11 AK019095	78887	1	similar to pir:T00322 - T00322 hypothetical protein KIAA0542 - human 62 %
11 AK004934	71774		Similar to APXL HUMAN Apical-like protein (APXL protein) humna 28 %
12 AK009086	69533		simitar to keratin associated protein 4.7 [31% Homo sapiens]
12 NM 023266	104348	Zfp120	zinc finger protein 120
12 AK018444	52504		expressed sequence Al448222
12 NM 013483	12231	Bintai	butyrophilin, subfamity 1, member A1
12 AK006472	74239		Rabe-interacting protein 2 [Mus musculus] 23.38 %
12 NM 008348	16154 Il10ra	U10ra	Interleukin-10 receptor alpha
12 AJ297743	30934	Tortb	torsin family 1, member 8
12 AK017387	71363		RIKEN cDNA 5430433J05 gene
12 NM 010351	14838	Gsc	george de consequence
12 AF358257			T46611 CL288 protein - rat (31 % R.norveglous)
12 NM 053168	94091	Trim11	tripartite motif protein trim11
12 AK015385	74673		RIKEN cDNA 4930451F05 gene
12 AK012063	67 18 1	Dullard	Dullard homolog (Xenopus laevis)
12 AK007540	69769		hypothetical protein FLJ23467 [Homo saplens] 93 %
12 NM_025554			
12 AF060246	22647 Zfp106	.fp 106	zinc finger protein 106
12 AK014490	73130		CA00_HUMAN Protein CGI-100 precursor (89% human)
12 AK013432	67608		Similar to nuclear pretamin A recognition factor, isoform a [84% Homo saplens]
12 AK011324			Mus musculus 10 days embryo whole body cDNA, RIKEN
		Eafl	Eaf1 protein
12 X72307	15234 H	Hof	hepatocyte growth factor
12 AK008409	67019 A	Actr6	ARP6 actin-related protein 6 homolog (yeast)
12 AF285585	30054 R	Rnf17	ring finger protein 17
12 AK008125	69831		adult male small intestine riken cdna ctone:2010005118
12 NM_019681	14299 F	Freq	frequenin homolog (Drosophila)
12 NM_019435	20130 N	Npts	Nuclear neuronal protein 15.8
12 NM_053192	94228 U	Ucc t	upregulated in colorectal cancer gene 1
12 NM 009125	<b>20239 S</b>	Sca2	spinocerebellar ataxia 2 homolog (human)
12 NM 008554	17173 A	Asd2	achaete-scute complex homolog-like 2 (Drosophila)
12 M12289	17885 M	Myh8	perinatal sketetal myosin heavy chain 3 end

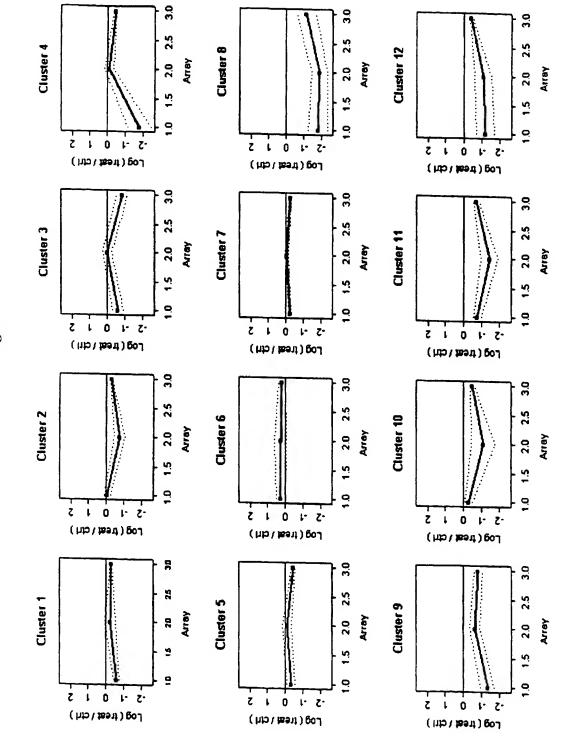


Figure 11-6

Description	Holocytochrome c synthetise	lectin, mannose-binding, 1	Eph neceptor A8	cytochrone P450, tamtly 2, suddamily d. polyocetide 26	per-hexamer repeal gene f	Nacety galactosammidase, applea	musicae specific gene (fromo sapients) 88 %.	acid beta glucosidase			much 10, submandibutar gland salivery much	Mus musculus actul male lestis cONA, RiKEN fulllength enriched library, cone;4933413N12	chapters give en unung protein tup-1			infrancease project	Under concerningers, yought sourcement e, o (mass mass ands) for the	muscle nicotinic acetylcholine receptor aighe aighe subunit aighe-subunit cholinergic	agun male cerebelarm raen cora cone; 15000 1407/	actives the expensive after 1 to deliver of the switch		vanin 3 PVETBERRANESS	CANTZPOSANOSZ protein jnomio sapransjos % / 100 88 ectonoleoside transcribate dibhosoholydiolose 5		TBC1 domain family, member 8	hypothetical protein FLJ20156 (Homo sapiens) 28 %	Corso protein [so a round saprems]	plecksim homology domath-containing, family A (phospholnositide binding specific) member 2	2210347C occaedin (Mus muscusus)	18 days embryo riken cana clone:1110005n14			beta-microsemmoprotein; beta-mhibin; prostade imitoin protein (Mus musculus)		OcuS histone acelytransferase; clone image:3491089		banscription factor like 4 toff4			T00335 hypothetical protein KIAA0564 • human (fragment)(69% human)	KIXEN COVA 1700128008 gene	camesponsive element binding protein 3 clone mgc:6348; ktp-1 and ktp-2 proteins 5end	ankyrin repeat domain 2 (stratch responsive muscle)	cancerheatis antigen 1; cancerfleatis antigen (Nomo sapiens) 36 %	menguc gregot ponni regulazion mine 2 1 minoribosofetal abbasomol peninim e 2 1	smaplobrava like 1	k+ voltage-gated channel subfamily s 2 kms2	adult male testis riken cona clone, 4933401b01		RIXEN COMA 4613427G06 cene	paired-like homoobox 2a	
į	Mccs	Lman	Ephas	Cvp2d28	Part	eden		g			Mucto	loff	Gzma	•		dund	200	Chma1	16 Apr	And 1		Vnn3	EntrodS		The 1d8			Plekha2	E	Maps			Msmb		GenSt		Tolle				Cmar	200	Ankrd2		Mag	Svori	Kcns2				Phox2a	
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Charles Access	1 NM_008190 1 AK008596	1 AK011495	6 MM 015707	1 NM 029562	1 NOM 01108D	1 NM 008669	1 NM 021493	1 NM 008094	1 AK005628 1 AK008518	1 AK006554	1 MM_008844	1 NM 019919	1 NM 010370	1 80006805	1 NM_025380	1 NM 008383	1 AK010062	I NM 007389	1 NM_025983	1 NM 009612	1 8C005624	1 NM_011979	1 NOM 007647	1 NM 053201	1 NOM_018775	1 AK010545	2 AK019361	2 NM 031257	2 AF151110	2 NM 015778	Z NSM_024277	2 AK012129	2 NOM_020597	2 AK005345	2 NM 020004	2 AK019788	2 NW 011550	2 AK014579	2 NUM_007863	2 AK004956	2 MM 020812	2 BC002094	2 MM_020033	2 NOM 025410	2 MM 008309	2 AJ133536	2 NIM 008436	2 AK018597	2 AK006389	2 AK014780	2 NAM_008887	2 AK018621

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12125	3. 3.		21420	26321	19047				13116	7633	Sign		17063	215705	:				17071	11687	10001	10821	12704	72183		52575		11567	20254	2077	225363		22262	12333	****	71941		69940	71519	66581	16023	00318	67267		69746		67072	19183	56470	12886			20613	78920	77634	80987	56469		21859		6781
Clories Access 2 AKO 1 185	2 NM 028119	2 NM_019578	2 NM_009335	2 MM_010210	2 NW 013636	Z MM 031173	2 AK016569	2 AK005730		2 AKO18281	2 AKD15427	2 AK015645	2 NM 010739	2 BC004091	2 AK018042	2 AK004913	2 BC004027	2 AK003556	2 NW 008530	Z NM 009660	2 030//3	2 NAV DOGUS	3 AF086824	3 AK011640	3 NAV 053186	3 AKD05047	3 NM 011246	3 AKU18083	New Post	3 NOV 023742	3 BC013717	3 AK003234	3 NM_009474	3 Nam 003802	3 104 009012	3 AK009937	3 NOV 010553	3 AK013041	3 AKOI 8458	3 AK003876	S ANA DAGGEO	3 ANTOLOS	3 NM 026063	3 AK012109	3 AK010555	3 AK008838	3 NOV 025850	COSSO MAN C	3 NOV 026446	3 NAV 009945	3 AK006262	3 AK014696	3 NM_011427	3 AK019713	3 AK018883	3 NOM 030729	3 NW 019683	3 NAV 010275	3 NOM_011595	3 AF302691	2 MM 015944

Description	guanidinoacetate methyltranslerase		spectrin beta 4		ASE				chondrolin 4-sufforansferase (Mus musculus) 47 %		small moudble cytotine a28 scya28		Mouse stock male diencephaton cONA. RDEN Ad Jength enrithed library, done 9330161L09 product hypothelical	suitotransterase-reated protein SULT-X1		cdc25a cdc25m3	entigen Identified by monocibral antibody MRC OX-2 recepto	myoloid/lymphoid or mixed lineage-leukemia translocation to homolog drosophila mili 10	KIAB (In) couled (Homo sections) 78 %	JCA913 anti-sigma cross-reacting protein fromotog I atoha precursor - 89% fuman	S-adenosytromocysteme hydrolase	RIKEN CDNA 1110020G09 gene	Mala mascalla (q. 11 daya emeryo wide body cures, rasen qui-emen emento normy, done 2010/1/12 product myonnessa Connessando		matha metalloprotestage 2 retigiz she after franse metalloprotestages 2 retigizes	auty zint, unger protein augin Ombren, zint, millen adult male testis riken coha clone, 4930570409		Chemokine (C/X-C modi) figand 12, 181182 Chokine - mouse 100 % AD16. Hi BJAN Dwilele AD018 (Periese CEL118) (A00001 90 %)	general banscription factor II A, 1				epsin 2	mitochondral cytochrome c oxidase subunit in cox exons 4 and 5; cox4	ubiquilin-conjugating enzyme E2M	Indicates inchrosoptatise (nucleosade biphosphate pyrophosphalase	caseholytic protease, ATP-dependent, protectic subunit homolog (E. coll)	10 11 days embryo risen cone cone 2810407111; cytochrome p450 701 cyp701	retinoic acid receptor garma, mrsi-cama-a	syntaxih binding protein 2	2-5 otgosdenylate synthelase 1G				prefoldin 2 pithn2	actual male formuse filtern catha chane;2310074e1\$		T 12488 Proofbetical contain DRF205840123 1 - Param 97 %	ribosomal protein mitochendrial I26 rpm28	and north annual for a	hepatoceflubr carchoma-easociated antigen 112 (55% Home sapiens)	108975 hypothetical protein CKFZp564F0522.1 - human (fragment) 49 % hypothetical postate El 110540 Herms saniens) 79 29 %	I house the second seco
8	Gant		Spub	12 13 14 15 15 15 15 15 15 15 15 15 15 15 15 15	1361	\$100310					82045 Call 2	*		Ser. S		Cdc25a	Mox2r	MIT 10			Ahcy		Gia5		Z drugg	Š		CKGIS	GIZa1				Epu2	S	Check First	eg	500	io o	. S. S.	Strbp2	Oasto				Ptdn2				Mrp(17	3	3		
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hypothetical protein A-211C8.1(87% human)
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Mus muscutis adul male small intestine CDNA, RIKEN full-length enriched library, clone;2010109K11 product:hypothetical protein
ubiquinin specific proteins 44
H-re-107. like protein 5 (Homo splens) 69 %
proteasome (prosome, macropain) subunit, alpha type 3
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RIKEN cDWA 2700007P21 gene
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Similar to PRP1 L. HUMAN Salvary protine-rich protein precursor (Clones CP3), CP4 and CP5) (Contatrs: Basic peptid 35 % human
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RIKEN-CDN4, 4221507707 gene
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Lucyd Porndog A (C. degans)
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YSC1 - KUMAN Putathe protein DJ7471423.2 (92 % human)
RINEK I Wild-Engh emiched (brany, chone:$4304094102
Cbptp300-interacting transactivator, with Glu/Asp-rtch carboxy-terminal domain, 4
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Similar to hypothetical protein Ft.J20546 (B2% Homo sapiens)
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RIKEN CDNA, 25 (10507), 03 (Mus musculus) 100 % of actory receptor 159
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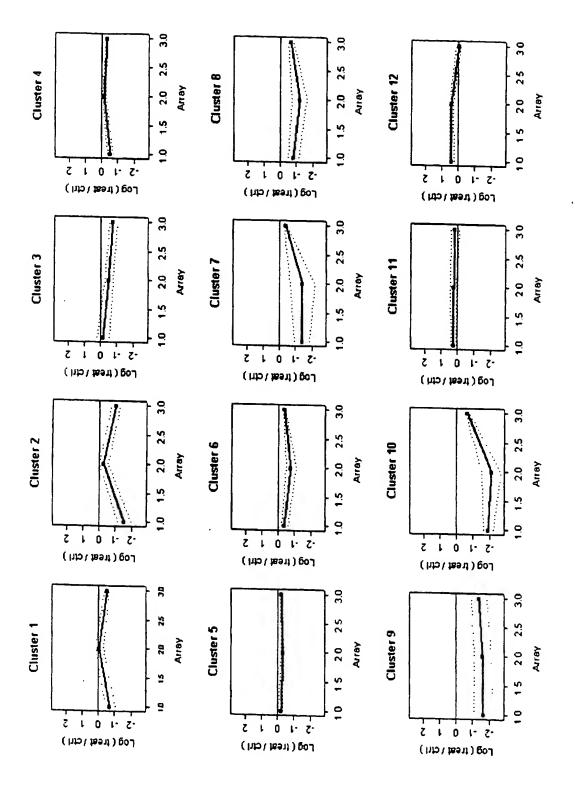
Description	M.muscutus ORF1 and ORF2 genes	Mus musculus 10 days embryo whole body cDNA, RIKEN full-length enriched library, clane;2610034B18 product;hypothelical pratein	Unyrold Mornone (eceptior-associated profesion fed in a second of a february school of the companies of the	independent of Graph Control of Graph Co	expressed sequence AI255456	ecotropic retrovirus receptor w i	phospholipase C, delta	Smitter to progesterore-from control (89% Homo exptens) DIFFER ATMA 23 FOR THE CONTROL OF THE CO	River Using 25 100 for the general and the second of the s	Eal orden	GTP binding protein 3	mucin 3, intestinal	T12515 hypothetical protein DKFZp4348103.1 • (28% human)	sphingosine phosphate lyase 1	cyclin-dependent kinase 5	zinc finger protein 316; kruppel-related zinc finger protein (Mus musoulus) 100 %	brain expressed myelocylomatosis oncogene	YOU domain, cases, a war source of the case of the cas	puspingue utern associated with approxpmingue eminated mindoonalins entit maje tests riken ochs done 489343705.	ras homolog gene family, member H	cytotoxic T-tymphocyte-associated protein 4	RIKEN CDNA 4930539A06 gene	nebutin-related anchoring protein	Solute carrier tampy 12 member 7	UUTT-humany makabengen nencasa prz (urzu-oox proten prz) (urzu-oox proten Tr) 49 % (	apolipomntein M	ribosomal protein mitochondrial 114 rpm114	gamma-aminobutyric acid gaba-a receptor subunit bata 2 gabrb2	pancreas specific transcription factor, 1a	RIKEN CDNA 28 05 10D 14 gene	nuolis (nuodespirate innee or moesy A-pype mour / A-positivativat noodes 1407 1407 1408 the second	hypotational process and the second s		RIKEN cDNA 0610009814 gene	Dullard homolog (Xenopus Isewis)	Perbox only profess 34, ref.NP_060413.1 - hypothetical protein FLJ20725 [Homo septens] 75 %	nings november 25 years 1875 MOLISE interescenting St. Da creates homeston (IFP 35) (100%, Mus musculus)		EST C77284	UDP-ducase detydrogenese	pelano I - Fernal VIV. 2013. I - Pelano I : Nuclear Curva, 20 I Vecca, 23 Ferna I I Marchina) I - U - N	TRUCKS TOCKET SWIND YOU WITH THE TOO TO THE TOO THE T	stander machindino protein homoloo drossonius staut; eduit mate liver riken odna chme:1300014004	atona 28-stalytransferase gd3 synthese putalive	GrpE-tke 2, mtiochondrtal	B hymphoid khrase	WAYT I MOUCH B SEPTEMBER DESIGNATION OF THE CONTROL	spylogistic see a ( North-Teodory, Catalogy of the see a see a X nov reneal commentation defective recent in Chimne handler relix 2	hypothetical protein R31449 3 - human (fragment) (89% human)	RIKEN CDNA 9130204G15 gene	Similar to myelokdrymphoid or mixed-tineago leukemia 2 (41% Human)	epithelial membrane protein - i	emylod bets (A4) precursor protein-binding, larraly A, member 3	onome trasseuke milochondrial ribosomal protein S18C; CGI-134 protein; milochondrial ribosomal protein S18-1 [Homo 78 %
8		;	Trap100	, 5		Skr7a1	Pic			Eafl	Gtobo3	Muc3		Sgpl	Caks	Z/p316	Bayc	10032	n n	Ach	Cita		Arab	Sic12a7		Abom	Mrbi2	Gobrb2	<b>E</b>		Mudi				Dultard	Fbx034			C77284	₽ Cod	N-4-3	28482	Starr	Slat8a	Grpet2	<b>8</b>	Wispt	Xeer?	1			Emp1	Appa3	Š
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1	3 X98456	3 AK011654	NM_011869	1 AK020739	1 AK012535	NIM_007513	NM_019676	MAK015/81	NM 028740	AK016628	AY029613	AF027131	AK014905	NM_009163	NM_007668	NM_017467	AK013780	MM_UUUUUU			Ī	-			BC003765				_		AKD04924	-	_		-	AK020018			_		•	AK010476	_	_	-		NM_018865	NM_0113/5	AK010963	AK020272	AK016775	NM_010128	NW_018758	AK004139
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Description	myociin	matriin 4	Protein tyrosine prospinates, non-receptor type 12	richt, or take-activating from I	dystican 1	p53 apoptosis effector related to Pmp22; p53 apoptosis-essociated target (Mus musculus) 100 %	oncogene-regulated cell acheston molecule orcam	comichan-like	high mobility group AT-hook 1	olfactory receptor 67	claudin 4	mucin 10, submandibutar gland salivary mucin [Mus musculus] 30 %	RIKEN cDNA 1500035H01 gene	bulyrophiin, subfamily 1, member A1	antigenic determinant of rec-A protein	calsequestrin 2	Mus musculus 10 days embryo whole body cDNA, RIKEN	boosecod dae	RIKEN CDNA 5430433J05 gene	upregulated in colorectal cancer gene 1	phosphoendpyrvale carboxykinase 1 cytosolic	Simula 10 Association control of the section of the	Sometime of programmer of the second	arcono certify triggerse o (cass v), pseudogene 1	Gracius Frenchiso median K. Frenchteins Peritain (Menna saniana) 99 Hindinadin bilassatina median K. Frenchteins Peritain (Menna saniana) 99	minore artisted rates those A	zinc (moer protein 108	transient receipt optential cation channel subfernity M. member 1	pre B-cel leukema transcription factor 3	hypothetical protein FL 123467 [Homo sapiens] 93 %		RIKEN cONA 2310016G11 gene	bone marrow stromal cell antigen 1	DNA segment, Chr 8, ERATO Doi 633, expressed	RIKEN CONA 6330583M11 gene	Similar to nuclear prefamin A recognition factor, isoform a [84% Homo sapiens]	odd Oznen-m homotog 2 (Drosophila)	A I P-binding casselle, sub-tarrity A (ABC 1), member 1	Spermin See	refregum 4 rrg4	and might pounded to the control of	mental company of the company of the master of the company of the	den interior manhena channel profess (	CAD HILLAN Price Control of California (85% human)	mannosidase 2. abha 82	RIKEN CDNA 4931431F19 gene	perinatai skeletai myosin heavy chain 3 end	hypothetical protein DKF2p434A1022 (Homo sapiens) 39.00 %	interferon response element binding factor 1	RIKEN CDNA C330006D17 gene	RIKEN CONA 4933406P04 gene	Smilar to APXL_HUMAN Aproachtike protein (APXL protein) humas 28 %	nepalocyte grown lactor	polassian voiagogated character desired sucharaly, order memory	nuclear factor of kappa upin polypetude gene emisercen in beceiss inminute, epsilon	ecopysecrated core, areasyse copyriously gain cent, sevaenary ore, ecopoem and usanon sequence of CEMPS shadeshed made and ecopysecrated of the copyright of t	Switz Smithau menuenance of chiomosomies z-ake i (yeas) mistosomal triaboseide transfer protein	The control of the co	histidine rich catchen binding protein	
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Access	C00010_MN	NM_013592	AKOOSOS1	NM 019825	NM_016696	NM_022032	NM_021339		NM_016660	NM_013619	NM_009903	AK007868	NM_023831	NM_013483	X58472	NM_009814	AK011324	NM 010351	AK017387	NM_053192	AKOOJESA	AKONASS	ACC4332	ACU15632 AF282291	NM OZESTA	x97052	AF060246	NM 018752	NM 016768	AK007540	AK007667	AK009387	NM_009763	AK017529	NM_024465	AK013432	NM_011856	NM_013454	NM_009214	MM_032002	NIA OORZAGO	AF282286	NM DRD450	AK014490	NM 008550	AK016497	M12289	AKD06911	NM 013714	AK021182	AKD16707	AK004934	K/230/	66C010_MN	OSSESSO MIN	M 025310	NM_008642	akma736	NM_010473	•
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Description	sp.095198 • KHL2, HUMAN Keich-like protein 2 (Actin-binding protein Marwen) 98 %	similar to ptr: T00322 - T00322 hypothetical protein KIAA0542 - human 62 %	transcriptional regulator, SIN3B (yeast)	RIKEN CDNA 2610208C17 gene	Nuclear neuronal protein 15.6	hypothetical protein MGC 15435 [Homo sapiens] 51 %	SRY-box containing gene 3	RIKEN CDNA 5730409E15 gene	keratin associated profein 13	phosphodlesterase 1C	fibrobast activation protein	CGI-94 protein	cholecystokinin A receptor	JC6547 high suffur protein B2E - rail 37 %	C-terminal PDZ domain ligand of neuronal nithe oxide synthase	similar to keratin associated protein 4.7 [31% Homo sapiens]	Ngfi-A binding protein 1	oliactory receptor 154	androgen recept	9030607L17Rix RIKEN cDNA 9030607L17 gene	Nii protein 2 (89% Homo sapiens)	BCL2-fike 12 (proline rich); Bct-2 like proline-rich protein 12 [Homo sapiens] 81 %	RIKEN cDNA 5730417817 gene	2-cell-stage, variable group, member 1; variable group of 2-cell-stage gene family	LSM3_HUMAN US snRNA-associated Sm-like protein LSm3 (MDS017) 100 %	nuclear factor of karpe light polypeptide gene enhancer in 8-cells inhibitor-like t	NADH dehydrogenase (ubiquirone) 1 beta subcomplex	mitochondrial ribosomal protein L4	alphe-globin transcription factor cp2		atp synthase h+ transporting mitochondral fifto complex subunit e atp5k; ifm-1 fifto-arpase	HSPC038 protein [Homo sapiens] 100 %	mdi10 deduced amino acid sequence of is horndogous to c. elegans putative dnaj protein z73102 b0035.14. horndog	RIXEN cDNA 1700006J14 gene	uncharacterized hypothalarrus protein HCDASE (77% Homo sapiens)		chemokine (C-C molif) ligand 4	Mus musculus adult male corpus stratum cDNA, KIKEN hall-ength entitined lugrary, clones. Co. condition product hypothetical hy	retiny Volgas, i hypothetical protein FLZUZZS [homo saptens] 65 %	policyrus receptor-related 3	complement component 1, q subcomponent, gamma polypeptide	Vacchina (Patient Kings)	expressed sequence AMA 6222	pusitive seven it answers that are spearang receptor purish	programmed cest deam to	Terri numana Procede pyrosionale-carooxysis populassi (3-oxoproprinassi) (17) opulasiny-populas 50 %	uand want of the control of the cont	AND	MMX non-receiptor types unage	MUS INDECEDED AND THE BEAUTION OF THE MICHENION OF THE BOARD OF THE BO	CERTIFICE BATTERING TRANSPORTER IN TRANSPORTER IN THE TRANSPORTER IN THE SECOND TRANSPORTER IN T	ACMS_MOUSE MUSICALISTIC ECCEPTOR MO (MILLS TIPOCITY) NO. #1		HSPC182 protein (98% Homo sapters)	GNAS (quanine nucleotide binding protein, alpha stimulating) complex focus	18 days embryo riken odna done:110002k21	proteotibid protein 2 (Mus musculus) 100 %	erythrobiasi membrane-associated pratein	microtubute-associated protein 18	chemokine (C-C motif) ligand 2	kinesin 9 kil9	opied growth tector receptor
8	KINIZ		Sus		Np15		Sox3	:	Krtap 13	Pdetc	Fap	Cg94	Cokar		Capon		Nab 1	Offr154	₹							NACHI	NdufbS	MrpA	Tcfcp2		Atp54		Onajb12			D6End772e	Š		!	E .	66.	VRZ		D-BLID	01808-1	!	C.		S T S	4			Cornel		Gras	3	Plp2	Ermap	Mtap 1b	S	5 5	je O
Locus	77113	78887	20467	72495	104130	72114	20675	70502	16699	18575	14089	67205	12425	96969	70729	69533	17936	27216	11835	71564	52633	75736	74737		67678	18038	66046	66163	21422	74282	11958	68036	56709	321010	52665	52440	20303	77478	66825	58998	12262	22669	2000	20,000	9750	70200	19799	99767	12169	CIRCOMO	12033	17444	20.5	16889	14683	66059	18824	27028	17755	20296	16578	72075
by Accras	AK020050			AK011897	_				Z :				_	_		-	Ž	_	_	AK018541	NM_023175	•	AK017569	•			_	NM 023167	_	_	_	-	_	•	_	Ž	_	-	•	NM_021496		-		מיסטרט אואי		11252U MM	_	-	MM_009/39	AK019053	DESCON MAIN	NM OTOBIS	NW 013657	AK019168	A.245739	NM 025327	NM 019755	NM_013848	NM_008634	NIM_011333	NM_010628	NM_031373
Clean	6	6	60	6	6	<b>o</b> n	<b>o</b> n 1	<b>6</b> 7 (	<b>6</b> 0 (	D) (	<b>5</b> 1 C	<b>6</b> 1	<b>o</b>	6	6	6	0	60	65	6	6	61	6	6	6	6	6	6	6	Ф	<b>0</b>	6	6	0	60 (	<b>о</b> :	2 9	2 9	2 9	2 9	2 9	2 9	2 9	2 9	2 9	2 \$	2:	= :	= :	= :	= :	: :	: =	=	=	=	=	=	=	= :	= :	=

Description DA.1 mDNA bluetien evaluis			peroxin 2, peroxisomal largeting signal 1 receptor-like; KIKEN cONA 1700016,008 gene [Mus musculus] 100 % T42372 mmballe ariandate kinase (FC 2.7.4.8). I membalas ariandate kinase (FC 2.7.4.8).	ORM Like 3 (S. cerevise)	RIKEN CDNA 1200011D03 gene	KIKEN CDNA 26105/38408 gene prof. Ownstand & ovidens 2 hanched chain	Nace Month Strategy Communication Communicat	arterial	neuro-oncological ventral antigen 1	selectin, platelet	Mus musculus adult male testis CDNA, RIKEN full-length enriched library, clone:4933427617 producthypothetical protein	ALEX3 protein	ubiquitin-conjugating enzyme E2 variant 1, isoform b; ONA-binding protein (77% Homo sapiens) shoomat contain 18	nouncedic inase (Homo saniens) 55 %	GNAS (guantine nucleotide binding protein, alpha stimulating) complex locus	RIKEN CDNA 2010004P11 gene	protein tyrosine phosphatase la-Zhela ptp an autoantigen in insulin-dependent diabetes mellitus; phosphatase-np ptp-np receptor	polaskum Gashine i modulado y lacido y como en cinamo e contrare e	chanozone 1 open reaming i ame 10 (nomo sapiens) 100 % historine enemine H 1	neuronein	steroid sensitive pene 1	small nuclear riboructeoprotein D3	RIKEN cONA 2410012H22 gene	sarcoglycan, alpha (dystrophin-associated glycoprotein)	hypothetical profein MGC2550 (84% human)	RIKEN cDNA 1500031J01 gene	methyltransferase Cy/19	RIKEN CUNA Z31003ZD18 gene	SAC ROTHOGOVY & CONTRAINING URANDONTHING POTENT U	Loyntra ase (trivia i i Urava de ecelo projectorale i NADH dehvitroeense (tripianinne) 1 albha subcomates 3	mesenchymal stem cell protein DSC54 (92.03% Homo sapiens)	transcription elongation regulator 1 (CA150)	isoprenylcysteine carboxyl methyltransferase	progestorae rembrarea binding protein (73% Homo Saptens)	dystronia modernia strass 815	G profein-coupled receptor 91	phosphoserine/threonine/tyroshie interaction protein	coiseniin-like protein	caterin alpha t	TAF11 RNA polymerase II, TATA box binding protein (TBP)-essociated factor	Cardony design from	KIKEN CAWA 120000MAD GENE	injector cer recentis acquerate a trephinosina, i reprintata abusantata interchalasa (Miss mascrinis) (M. 60. embientatas i abbestata abusantatasa 1 - embinosina, i abusantata abusantatasa (Miss mascrinis) (M. 60.	Janus kinase 2	RIKEN CONA 1200017A24 gene	MAs musculus 15 days embryo embryonic body below displaragm cDNA, RINEN Adl-length enriched library, close;8230401417 product:hypothetical protein, full insert sequence	truncated SON protein [Mus musculus] 34 %	KLKC HUMAN Kalikrem 12 precursor (Kalikrein-like protein 5) (KLK-LS) 70 %	South Carrier tarmy 30 (zur entsporter) menner b	myelood eeconopie, kirsa hijepigaaloon jajerkeakoo genee k kirooleen keksenan sendain (1178 Ribono santama 69 k.	unterent educin ruder ent process control septembly on an impantite motif protein 34
Oene	Case (A	Nassio One	rest	Ormdia		Acres 2	Nack	Art	Novat	Sep		Alex3	8	2	Gnas		Ptprn2	E C	HTP	N	Ssg1	Snrpd3		Sgca			Cyrig	į	o de	Nduta3		Tcerg1	Ē	Ē	Om15	60G	Shya	Calp	Catne	Teff	3	1	S	Jako					2000	76.m	Trim34
Locus	73246	05767	52357	66612	71724	61719	56174	11876	18134	20344	74466	71703	36887 26961	69168	14683	67889	19278	60018	15465	56183	87896	67332	69747	20391	68520	67529	57344	74182	20420	16099	71373	56070	57295	10804	13400	84112	56291	80334	12385	68776	12613	4106	81575	16452	74146	77550	66239	69511	210148	17337	94094
Chotche Access	11 AK005472	THE COLUMN		11 NM 025681	11 NM_023617	11 NM 053115	11 NM 019542	11 NM 009711	11 AF232828	11 NM 011347	11 AK016951	11 80011101	11 NM 012053	11 AK007531	11 NIM 010309	11 NIA_026434	11 UR2439	11 AK/07751	11 NM 008285	11 NM 019515	11 AK011256	11 NM_026095	11 AK010472	11 NM_009161	11 AK003661	11 NM_026218			12 AKO12818		12 AK017282	_						12 NM 030265			12 KM COSBBS			12 NM 008413	12 NM_028785			12 AK009217	12 BC005/53	12 AKM2852/	12 NM_030684





Cluster Access	Locus	Gerra	Description
1 NM_020025	26878	B3gall2	udp-gal:belagicnac bela 13-galactosyftransferase potypeptide b3galt2
1 NM 011408	!		
NM 016880	53617	Kd1-24	hair keralin acidic 5 ha5
1 AK014408	74315		patched related protein translocated in renal cancer [28.14% Homo sapiens]
1 NM_019774	26399	Akap8	A kinase (PRKA) anchor protein 8
1 RC005625			
1 NM 030750	81535	Sapp1	sphingosine-1-phosphate phosphatase 1; sphingosine-1-phosphate phosphatase (Mus musculus) 100 %
1 AF 264049			
1 NM_010340	14765	Gpr50	G-protein-coupled receptor 50
1 AK016419			
1 AF285178	16351	đ	actin-binding protein mipp
1 AK017880			
1 AK019493	75735	Pank 1	pantothenate kinase 1
1 NM_016788	51789	Tnk2	tyrosine kinase, non-receptor, 2
1 NM_007928	13728	Mark2	MAP/microtubule affinity-regulating kinasa 2
1 NM_019544	56184	Msgn1	pMesogenin 1; mesogenin (Mus musculus) 100 %
1 AK017242			
1 BC003298			
1 BC010539			
1 NM_026495	67991		RIKEN cDNA 06 10020102 gene
1 NM_008697	18080	ninein	ninein
1 AY028606	114566	Krt2-20	ref:NP_149022.2 - keratin, hair, basic, 2; hard keratin, type II, 2 [Homo sapiens] 85 %
1 NM 010063	13426	Dncic1	dynein cytoplasmic intermediate chain 1
1 NM 026570	64050	Gas41	glioma-amplified sequence-41
1 NM 009212	20589	1ghmbp2	immunoglobulin mu binding protein ighmbp2
1 NM 011427	20613	Snail	snail horrolog drosophila
1 NM 008393	16373	tx3	iroquois homeabox protein 3
1 AK009177	70088		Similar to hypothetical protein FLJ11730 [97% Homo sapiens]
1 AK010201	69687		CGI-127: yippee protein [100% Human]
1 AK002546	66058		hepatocellular carchoma-associated antigen 112 [55% Homo sapiens]
1 NM_010080	13517	Osbb	dentin siatophosphoprotein
1 AK018594	68145		ref:NP_061875.1 - ETAA16 protein [Homo sapiens] 48 % /
1 AK002747	71685		hypothetical protein FLJ12691 [81% Homo sapiens]
1 AK020207			
1 NM_011519	50969	Sdc1	syndecan f
1 AJ293897			
1 NM_007619	12402	₹	c-cbl proto-oncogene
1 AK004206	67282		AD16_HUMAN Protein AD-016 (Protein CGI-116) (x0009) 90 % /
1 NM_013696	22045	Trific	thyrotrophin-releasing hormone receptor trh-r; thyrotropin releasing trhr
1 NM_012043	26968	isi Si	immunoglobulin superfamily containing leucine-rich repeal
1 NM 033374			
1 NM 007706			
1 AJ237585	54392	Hcapg	chromosoma condensation protein G
1 AK014783	73914	lrak3	interleukin-1 receptor-associated kinase M (73% human)
1 NM 011017	18408	Slc25a15	solute carrier family 25 mitochondrial carrier omithine transporter member 15 slc25a15
1 NM 018827			
1 NM_011665			
1 AK009836			
1 AK017809			
1 NM 026580	68149	:	ubiquitir-specific protease otubain Z (94%)
1 AJ133536	20955	Sybii	synaptobrevin like 1

Description	flavin containing monooxygenase 1	OnaJ (Hsp40) homolog, subfamily B, member 1 FXVD domain-containting fon transport regulator 1	dentatorubrai pallidoluysian atrophy	mesoderm posterior 2 mesp2 odc25a odc25m3	mel-1a melatonin receptor	RAB38, member RAS oncogene family		acyt-Coenzyme A oxidase 3, pristanoyt; pristanoyt-CoA oxidase [Mus musculus] 31 %		calcium channel, voltage-dependent, beta 3 subunit	747230 humathatian pantain DKCZA4248002 1. humana (francount) 95 tk	amyloid beta protein precursor beta-amyloid; hippocampal	hypothetical protein FLJ22344 [85.6% Homo sapiens]	ubiquitin specific protease 29	angiopoietin	myomesin 2 myom2	Cystatin B	Contract in your cooper (A, rearis CTA) lind	2113200G ribosomal protein S10 (98% human)	Mouse 10, 11 days embryo whole body cDNA, RIKEN full-length enriched library, clone:2810417J12 product:hypothetical		adult male testis riken cdna clone:1700066j24	mpv17	adult male lestis nken cdna clone: 1700023a16		preprosomatostatin	done mgc.7522			sulfotransferase sulf1b1 washu-hhmi est genbank accession number aa267283; dopa/lyrosine	taminin atpha-chain tama5 basal tamina/basement membrane component	ropporin loc81026		
Gene	Fmot	Dnajb1 Fxyd1	Orpla	Mesp2 Cdc25a	Minrla	Rab3b				Cacub3		App		Usp29	Agpt	Myom2	\$ 2	3					Mpv17			Smst				Sult1b1	Lama5	Rppn		
Locus	14261	81489 56188	13498	17293 12530	17773	69908		74121		12297	76464	11820	78771	57775	11600	17930	13014	01710	67097	76917		76992	17527	69371		20604	66310			56362	16776	76378		
Cluster Access	2 NM_010231 2 NM_009358 2 AK009532 2 AK003234	2 NM_018808 2 NM_019503	2 NM_007881	2 NM_008589 2 NM_007658	2 NM_023805 2 NM_008639	2 BC011109 2 NM_023537	2 AK007262 2 NM 023644		2 NM_025592 2 AK015645	2 NM_007581	2 NM_033327	2 NM 007471	2 AK013379	2 NM_021323	2 NM_009640	2 NM_008664	2 NM 007793	2 NM 023850	2 NM_025963	2 AK013108	2 NM_008621	3 AK006904	3 NM_008622	3 AK006257	3 AK011730		3 BC002240	3 0/0139	3 NM 025331	3 NM 019878	3 U37501	3 NM_030744	3 AK014595	3 AK013037

Cluster Access	Locus	Gene	Description
3 AVA 4234	15381	Harpc	heterogeneous nuclear nbonucleoprotein c hnrnpc
3 NM COOSOA	222	0	PULZ, MUUSE Kettövints-realied PUL potytatiem (Contains: Reverse transcriptase : Endonuclease)(32 % M.muscutus)
3 AB000121	n † †	Simily.	cholinei gic receptor incollinic gantina polypeptide ching; muscle acetycholine gamma-subunit
3 AK011555			
3 NM_054097	117150	pip5k2c	phosphatidylinositol-4-phosphate 5-kinase, type II, gamma
3 NM_013541			
3 AK014579			
3 AK003384			
3 NM_025937			
4 AK014438	67111		Similar to PLT_HUMAN Protein PLT 81%
4 NM_028605	73682		A44492 probable DNA/RNA-binding protein - rat 55 %
4 AK018269	76877	Rab36	RAB36, member RAS oncogene family
4 AK012416	67088	Tp120b	TBP-interacting protein b
4 NM_021481	58866	Treh	trehalase (brush-border membrane glycoprotein)
4 KM 012008	00000	4	in the state of th
4 NA 015575	2220	CWXOL	- Cox and we-to domain protein 5
4 NM 020271	67078	Cours	Acquair ANCh 4103
4 AK021365	77998	Criffe	ripponitrional protonitri in 100 - 110 million - 100 million in ordere bladische britaine beste 27 il -90 hadres biadisce sected
4 NM 022993	55107	5	processing the control of the contro
4 AKOOROGO	60870	2	TIPS OF COUNTY STORMS TO CONTROL OF COUNTY OF
4 NM 022315	64074	Smort	SDARC relation modular relation bindion 2
4 NM 008657	1787R	Md6	md.B
4 AKO16704	24430	myio	Tributan 2 (2018) 14 months of the contractions of the contraction
4 ANGIOVO:	14430		gene trap locus 3 (27% m.musculus)
A NIM DIBYRI	50273	Pex14	protectional blogeresis later 14. FEXE_MOUSE Peroxisonal membrane protein PEX14 (Peroxin-14)PEX1 100 %
5 NIM 031247	1,667	P or a	FAREN COUNTY OF VALORISMS
5 BC004803	90409	Ž	HITHING GOSOCIALED INCIRCULOR IN ME
5 NM 025397	66172		RIKEN JONA 111000 Mile misser this 100 %
5 AKO21056	77314		CATALA COLON CONTROL MAN INCAMAGE TO A CATALA COLON CATALA COLON CATALA COLON CATALA C
5 NM 025435			
5 AK009247			
5 BC003289			
5 NM 007859			
5 AB049954			
5 AK010002	69668		RIKEN CDNA 2310061109 gene
5 NM 053194	101489	AI114950	synembryn
S NM_008075	14408	Gabrr1	gamma-aminobutyric acid receptor rho 1 subunit ionotropic gaba
5 NM_023380	67742	Samsn1	SSNI_MOUSE SAM-domain protein SAMSN-1 (SAM domein, SH3 domain and nuclear localisation prote 100 %
5 AK015616	74953		JC4194 lanosterol synthase (EC 5.4.99.7) • human 84 %
5 NM_008418	16491	Kcna3	polassium voltage-gated channel, shaker-retated subfamily, member 3
5 NM 022305	14595	B4galt1	UDP-Gai:belaGlcNAc beta 1,4- galactosytransferase, polypeptide 1
S NM_031385		,	
5 NM_053011	94217	Lrptb	low density lipoprotein-related protein 18 (deleted in tumors)
5 AK018759	66943		hypothetical protein FLJ22378 (81% human)
5 NM_016689	11828	Aqp3	aquaporin
5 NM 007442	11695	Alx4	aristaless 4
5 NM_019942	56526	38961	septin 6
5 NM_010582	16425	Itih2	inter-alpha trypsin Inhibitor, heavy chain 2
5 AK018624	71581		RIKEN CDNA 9130015AZ1 gene

5 NM 026415 5 NM 02635 5 NM 02635 5 A13825 5 A838257 5 NM 028717 5 AK015359 5 NM 02856 5 NM 030690 6 NM 038311 6 NM 028311 6 NM 028311 6 NM 028311 6 NM 028311 6 NM 028311	67769 12572 70967 74018 54563 14912 72561 72561 72112 72112 72112	Cdk7 Als2 Nup210 Nw6-2 Nkx6-2 Terf1 Terf1	cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating kinase)  T466 11 CL2BB protein - rat (31 % R. norvegicus) amyotrophic lateral sclerosis 2 (juvenile) homolog (human)  nuclear pore membrane glycoprotein 210 poin 210  NK6 transcription factor related. Iocus 2  DKF2P564G013 protein; novel retinal pigment epithelial gene; novel retinal pigment epithelial gene 84 % hypothetical protein FLJ2 1168 [76% Homo sapiens]  Similar to protection of telomenes 1: DKF2P56BD211 protein [71% Homo sapiens]  PKC-dependent PP1 inhibitory protein subunit 14 [Mus musculus] 100 % telomoric repeat binding factor 1  defense/immunity protein activity, indoleamine-pymole 2.3-dloxygenase activity
6 AKO10800 6 NM_010800 6 NM_010802 6 NM_01705 6 NM_01705 6 NM_01920 6 NM_01920 6 AKO15374 6 AKO14334 6 NM_020578	76797 18192 63986 17984 22367 76571 76571 2633 20630 70887	Nsconi Gmfg Ndn Vrk1 Abcg2 Smp1c	apprayment conscription record of the seas I histocompatibility antigen H-2 M3 alpha chain precursor - mouse 62% non-selective callon channel 1 glia maturation factor, gamma necdin vaccinia related kinase 1 map kinase phosphatase-like protein MK-STYX [Homo sapiens] 64 % ATP-binding cassette, sub-family G (WHITE), member 2 hypothetical protein FLJ23356 [Homo sapiens] 81 U1 small nuclear ribonucleoprotein 1C Mus musculus actul male testis cDNA, RIKEN full-length enriched library, come:4921520P21 product.hypothetical Microbodies C-terminal targeting signal containing 3 EH-domain containing 3
6 NM_01154 6 AK012224 6 NM_01352 6 AK016890 6 AF052942 6 NM_033561 6 NM_003561 6 NM_008325 7 AB041649 7 AK012378	19054 76795 20303 7 1099 13143 76559 17386 21390	Ppp2r3a Ccl4 Dapk2 Mmp15 Tbxa2r	protein phosphatase 2 (formerly 2A), regulatory subunit B, alpha vascular Rab-GAP/TBC-containing: BUB2-tike protein 1 [Mus musculus] 48.54 % chemokine (C-C motif) ligand 4 serine(Ihreonien kinase FKSG81 [Homo sapiens] 44 % death-associated kinase 2 Similar to hypothetical protein FLJ10242 [95% Homo sapiens] matrix metalloproteinase 15 thromboxane a2 receptor
7 AK01779 7 AK018940 7 NM_054039 7 NM_055798 7 AK015001 7 NM_011248 7 NM_011248 7 NM_013959 7 AK019494 7 NM_033567 7 NM_033567	70163 20371 66733 19649 13605 78261 94047 21343	Foxp3 Rbig1 Ecl2 Cecr6	Mouse 8 day embryo whole body cDNA, RIKEN full-length emiched library, clone:57305191.10 product:myristoylated alanine rich protein kinase C substrate Similar to A43932 mucin 2 precursor, intestinat - human (fragments) 27 %  EXP3_MOUSE Forkhead box protein P3 (Scurfin) 100 %  Similar to potassium voltage-gated channel, subfamily G, member 1; potassium channel KH2 [78% Homo sapiens] retinoblastoma inhibiting gene 1 ect2 oncogene  Mouse 0 day neonate skin cDNA, RIKEN full-length enriched library, chone:4632413E21:weakly similar to rat phosphotipose 8 cat eye syndrome chromosome region, candidate 6 homolog (fruman)  TAF6 RNA polymerase II, TATA box binding protein (TBP)-associated factor

Description frizzled homolog 4 drosophila fzd4 RIKEN cFNA 3732407/237 send		io i i delys entaryo inten cana cione; 28 i 040 / i 1 i, cytochrome p450 7b1 cyp7b1 C-ret proto-oncogene		zinc finger protein 162 Elevative armuth footbar 33				cyclin-dependent kinase 2 [Mus musculus] 100 %	phospholipase A2, group IID	hypothetical protein MGC 10771 [78% Homo sapiens]	RIKEN CONA 201001E1 1 gene Marco actif mala todia PMA Divental fronts actif to active	moosa avan maa kasas cura,, mten ulmengur emitaka ilorary, donesasuozorzi produdinypunelidal proferi (vrosine phosphalase recentor brae O	similar to hypothetical protein FLJ10008 (Homo sapiens		Mouse adult male testis cDNA. RIKEN full-tenoth enriched library, chone:4821530D09 omduret-humbheir-al	erythrocyte protein band 4.1-like 3	e adult male kidney riken cdna clone:0610009o09			B-box and SPRY domain containing	SINCE SECRET OF CATE OF THE SECRET OF THE SE	CINENT TIME 4501 4517 18 GENE	SIY-TOOK COMMANDED SIYOU RIKEN FONA 4970F58 124 nama	paired box gene 2	9030409E16RIK RIKEN cDNA 9030409E16 gene	protactin-tike protein C 2	RIKEN CDNA 1700013004 gene	RIKEN CDNA 4930473B18 gene	neuroligin 1	Nomeodornain prolein cix homeodox	pulative seven i larismentane spanning receptor puna-g RIKEN cENA 493-507P07 nena	milochordrial solule carrier protein	interferon induced transmembrane protein 4 like	putative nuclear protein ORF1-FL49 [Homo sapiens] 89.42	Mouse adult male testis cONA. RIKEN full-length enriched library, clone:1700122C07 product:serine/threonine khase 33.	potassium large conductance calcium-activated channel, subfamily M, alpha member 1	C-type (calcium dependent, carbohydrate recognition domain) lectin, superfamity member 6	aldo-kelo reduciase Jamily 1, member C18	keralin associaled protein 13	TDA consistent locus	Transguater Organists of State	Coll annual remaining with Fight and dampin Home canions	gene for odorant receptor A16	
Gene Fzd4	Pax5	Ret	Rnasep2	Zíp162 Enf22	rgicz Laste 12	Zlp354c	Arts1	Cdk2	Pla2g2d			Ptoro				Epb4, 113	D15Wsu77e		1	Bspry	Smck	500	YOS	Pax2		Prtpc2		;	Zign 1	خ د	r F F F F F F F F F F F F F F F F F F F	Mscp	Mik			Kcnma1	Clecsf6	Akr1c18	Клар13	Tood	- Clord	5	Ora16	
Locus 14366 74014	18507	19713	54364	22668 67112	56072	30944	80898	12566	17782	74577	72045	19277			77058	13823	110961			021261	20591	0807	75342	18504	66817	67310	75502	75803	192167	12951	70821	67712	74482	99099		16531	26888	105349	16699	21002	72068	68567	57272	
Cluster Access 7 NM_008055 7 AK014404	7 NM 008782	7 NM_009050	7 AK004137	7 AKO08922	7 NM 019516	7 NM 013922	7 NM_030711	7 NM_016756	7 NM_011109	8 AK014667	8 AK008003 8 AK015893	8 NM 011216	8 BC002230	8 AK016572 8 AK009020	8 AK014983	8 NM_013813	8 NM_033073	8 NM_008672	8 AK00/269	8 84276690	8 AF12/245	8 NM 04446	8 AKO15148	9 X55781	9 AK018496	9 AF158744	9 AK005970	9 AK015565	9 BC005523	077700 MN 6	9 AK014840	9 AK019700	9 AK017110	9 AK002512	9 AK007235	9 NM_010610	9 NM_011999	9 AB059565	9 NM_010671	0 MM 011626	9 AK011231			

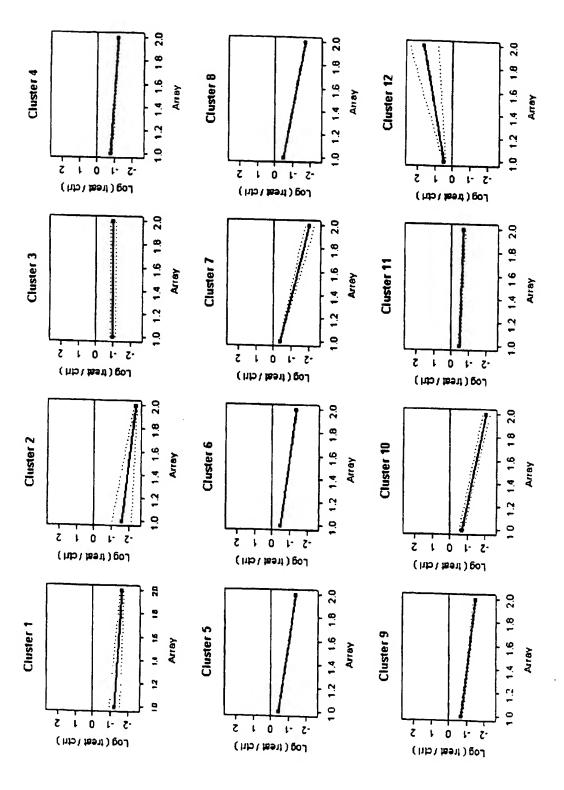
	Description	KIAA10/4 prolein [Homo sapiens] 45.33 %	cytochrome P450, family 4, subfamily f, polypeptide 16	adult male testis riken odna clone:4922502b01	transcription factor ap-2 beta tcfap2b	10 days neonate head riken cdna clone: 5530401a14	RIKEN cDNA 4921509822 gene	ATPase, H+ transporting, V1 subunit C, isoform 2	RIKEN cDNA 1700080G18 gene	CDS2 antigen	e2f transcription factor 5 clone mac:6043: e2f-5 protein	hypothetical protein MGC26988 (38% human)	RIKEN CDNA 2310016N21 (Mus musculus) 100 %	prolactin-like protein A	RIKEN CDNA 4930504H06 gene	male-specific lethal-3 homolog 1 (Drosophila)	tubertn-like protein 1	RIKEN cDNA 2310046K23 gene	adult male tongue riken cdna clone:2310035/15	phospholipase C, zeta 1	neuropaptide y receptor y1	olfatory receptor mor17-1	RIKEN cDNA 1110019L22 gene	RIKEN cDNA 4921517D21 gene	fragile histidine triad gene	lymphotoxin A	syntrophin, acidic 1	adult male testis riken cdna clone:4930578005	RIKEN cDNA 3200001D21 gene	Mus musculus mRNA for CN 8 scFv, complete cds	phospholamban [Mus musculus] 100 %	RIKEN CDNA 1700006C06 [Mus musculus] 100 %	sarcoglycan, alpha (dystrophin-associated glycoprotein)	plectin 1, intermediate filament binding protein, 500kD [Homo sapiens] 25.74 %	solute carrier family 21 (organic anion transporter), member 13	phosphalidylinositol glycan, class A	RIKEN cDNA 9530004P13 gene	olíactory receptor 64	seaeled frizzled-related sequence protein 5	FYVE and colled-coil domain containing 1	FK506 binding protein 11	Akr1c6	seminal veside antigen-like 2; SLP in the mammary gland; SVA-like protein in the mammary gland [Mouse 100 %	membrane protein, palmitoylated 5 (MAGUK p55 subfamily member 5)	histocompatibility 2, M region focus 2	vav2 oncogene	hypothetical protein FLJ10569 [100% human]	golgi associated, gamma adaptln ear containing, ARF binding protein 2	RIKEN CDNA 2310032D16 gene	RIKEN CDNA 4933436E20 [Mus musculus] 100 %	hypothetical protein FLJ20473 [Homo sapiens] 69 %	sulfotransferase, hydroxysteroid preferring 2	Iransmembrane 4 superlamily member 1
•	Gene		Cyp4116		Tcfap2b			Atp6v1c2		Cd52	E215			Pripa		Msl31	Tulip1			Picz 1	Npy 1r	mor17-1			F.	c <sub>1</sub>	Snta1	Hist4			돈		Sgca		Slc21a13	piga		Offr64	Strp5	Fyco1	Fkbp11	Akr1c6	Sval2	Mpp5	H2-M2	Vav2		Gga2				Sth2	Tm4sf1
	5000	70,02	70101	70954	21419	71388	70859	68775	73533	23833	13559	70897	75581	19110	75040	17692	56784	76968	69585	114875	18166	209102	68530	67722	14198	16992	20648	15269	71818		18821	66320	20391	74902	28254	18700	77363	18365	54612	17281	66120	83702	84543	56217	14990	22325	70885	74105	74182	71202	71699	20865	17112
	CHURCH ACCESS	STATION A	9 NM 024442	9 AK015021	9 NM_009334	9 AK017430	9 AK014850	9 BC003810	9 AK006956	9 NM_013706	9 NM_007892	9 AK014851	9 NM_023784	9 NM 011165	9 AK015697	9 NM_010832	9 AK009111	9 AK009853	10 AK009636	10 NM_054066	10 NM_010934	10 AF133300	10 AK003826	10 NM_026338	10 NM_010210	10 NM_010735	10 NM 009228	10 AK016310	10 AK014278	10 AB036341	10 NM 023129	10 NM_025486	10 NM_009161	10 AK015166	10 NM_023718	10 D26047	10 AK020544	10 NM_013616	10 NM_018780	10 AK013060	10 NM 024169	11 NM_030611	11 NM_032542	11 NM_019579	11 M26156			11 AK004632	11 AK009563	11 NM_027719	11 BC011108	11 NM_009286	11 NM_008536

Description	RIKEN CONA 4833224K13 Dene	DIFFER AND STROKED STROKE	NINEW COUNTY STOOT DEAD UP TO THE PROPERTY OF	KIKEN CDNA 4921525L17 gene	RIKEN cDNA 8430319H21 gene	days embryo niken odna clone:5730521f22	epireaulin	outstive N-acetylransferase Camello 4: RIKEN cONA 06:100370-16 neuro l'Aus musculus) 100 %	CLDS MOUSE CIANTINA, 2.3%	DIVERSION OF A CONTRACT OF A C	KINEN CUNA 1810014F TU GENE	day neonate skin riken cdna done:4632415i09	exportin 5	Z208_HUMAN Zinc finger protein 208 53 %	cystatin 8 (cystatin-related epididymal spermatogenic)	immunoqibbulin superfamily member iqsf6	Mus musculus Trif receptor-associated (actor a (Trails) overs, partial sequence; amplionless precursor profeso				recentor tyrosine kinsse-like omban nort - ROR1 MOUSE Tymolae-roniein kinsse fransmembrane nerestor ROD t	hynothetical profess FI LINES (35%, hymnos)	Knimelike (adne 1 endhede)	Properties and Expendence		Comparing manufactured and an advantage (A)	o koreji kondijan ja de koreji kondijan ja de koreji kondijan ja de koreji kondijan ja de koreji kondijan ja d	checochode Checken 9 Hanne engineer (CD D)	grycopitain C. Isorom Z (nomo sapiens) od %	embryonic lethal - Es2 protein. DNA segment, Chr 16, human D22S1269E, expressed (Mus muscutus) 100 %	galadosidase, alpha	offatory receptor mor 17-1	contactin associated protein 4	simitar to RtKEN cDNA 1810006A16 gene (Homo sapiens) 84 %	lysophospholipase 1	c-fos cellular homolog to viral oncogene c-fos protein	RIKEN cDNA 4921508M14 gene	T00335 hypothetical protein KIAA0564 - human (fragment)(89% human)		10 11 days embryo riken cdna ckone;2810431n19; ctone image;3498575		RIKEN cDNA 1700026G02 gene		topoisomerase (DNA) III alpha	neuromedin	RIKEN cDNA 4930557821 gene		859254 mysoin heavy chain 12, splice form2 - human 29 %	uterine-specific proline-rich acidic protein	hematological and neurological expressed sequence 1	zinc finger protein mkr5 3 end		
Gene						Sach	Fred	Cm.				Hsp25	Xpo5		Cst8	gstj	ò				Ror1		KII	Duct	i A		OEIDO			Eszel	eg S	mor17-1	Cntnap4		Lypla1	Fos				Trim29				Top3a	N E N				25	Ī	Zfp28		
Locus	75770	60563	2000	8160/	77849	24051	13874	68396	73347	60064	40000	15507	72322	170763	13012	80719					26563	75137	16596	24018	200	07000	76008	71687	2007	989/7	11605	209102	170571	73608	18777	14281	70844	71758		72169		75547		21975	56183	75308		74717	22264	15374	22690		
Cluster Access	11 AK014760	11 AK009351	00000000	11 AKU14968	11 AK021026	11 AK017781	11 NM 007950	11 NM 023455	11 AK006680	12 AKOO2494	10101010 7:	12 AK019498	12 AK012248	12 AF242377	12 AF090691	12 NM_030691	12. AF320615	12 AK007707	12 AK005311	12 AK014062	12 NM 013845	12 AK015972	12 NM 010635	12 NM 011884	12 NM 008085	12 MM 030726	12 AKOO7281	12 AKOO2769	12 ANOC103	12 NM 022408	12 NM_013463	12 AF 133300	12 AF333770	12 AK007346	12 NM_008866	12 NM_010234	12 AK014845	12 AK004956	12 NM_025675	12 BC006699	12 AK015864	12 AK006382	12 AK014485	12 NM_009410	12 NM_019515	12 AK016159	12 NM_009792	12 AK015780	12 NM_009475	12 NM_008258	12 M36516	12 AK014391	

Figure 14-8

ne Description	gn regucalcin	expressed sequence AI256456	•		Ū	JC6547 high sulfur protein B2E - rat 37 %	Ŭ	p fibroblast activation protein	_		.2 vaccinia related kinase 2	_	5 endotheltal differentiation, sphingolipid G-protein-coupled receptor, 5			ab methylmatonic aciduria (cobalamin deficiency) type B homotog (human)	_	51 G two S phase expressed protein 1		M.musculus ORF1 and ORF2 genes		hypothet, ical protein FLJ10694; hypothetical protein FLJ10889 [87% Homo sapiens]	Ī	RIKEN cDNA 0610042E07	12 protein tyrosine phosphatase, non-receptor type 12	hypothetical protein, MNCb-2622	YYY transcription factor	2 keratin complex 1, acidic, gene 2	c1 chloride channel K1	Rab6-interacting protein 2 [Mus musculus] 23.36 %	11 butyrophilin, subfamily 1, member A1	antigenic determinant of rec-A protein				adult male urinary bladder riken cdna clone:9530081k03	_	DNA segment, Chr 1, ERATO Doi 396, expressed - RIKEN cDNA 2610307121; hypothetical MNCb-4273 (Mouse) 100 %	tissue inhibitor of metalloproteinase 1	lymphocyte-activation gene 3
Gene	Rgn		Aqp2	8rms1	SS		Casq2	Fap		Pdefc	Vrk2	Myo1b	Edgs	C149		Mmab	Col4a3	Gise1	Rad541		Ē		Epcs3		Ptpn12		Yy1	Krt1-2	Clcrk1		Btn1a1	X Si	Mga	)	Nab1		Myg1		Timp	Lag3
Locus	19733	101513	11827	107392	14836	96969	12373	14089	69578	18575	69922	17912	14739	12262	52504	77697	12828	29870	19363		214162	101437	56092	67441	19248	58227	22632	16670	12733	74239	12231	16588	29808	72125	17936	78558	60315	52477	21857	16768
Access	1 NM_009060	1 AK012535	1 NM_009699	1 AF233580	2 NM_010351	2 NM_027170	2 NM_009814	3 NM 007986	3 AK009387	3 NM 011054	3 AK012664	3 NM_010863	4 AF108020	4 NM_007574	4 AK018444	4 AK020286	5 NM_007734	5 NM_013882	6 NM_009014	6 X98456	6 L17069	7 AK014260	7 NM_025310	8 NM_026158	8 NM_011203	9 NM_021416	9 NM 009537	9 NM_010665	9 NM 024412	10 AK006472	10 NM_013483	11 X58472	11 NM 013720	11 AK011185	11 NM 008667	12 AK020653	12 AK017955	12 NM_021421	12 NM 011593	12 NM_008479
Cluster																																								





	ma helicase a doxe and drosophila mie aduli male lestis riken odna clone:1700123:008
ž 8	metal response element bridging transcription factor 2 CDKZ (cyclin-dependent kinase 2)-associated protein 1 n-wasp binding protein wish substance p receptor
Mus muscuhu gatactosidase Ellis van Crew NK1 irenscrip sneil homolog sneil homolog cadherin cah6	Mus musculus RIKEN cDNA 1600010003 gene, mRNA (cDNA ctone MGC:7480 IMAGE:3490700) galactositase, aiphe Eliis van Greveld gene homolog (turnan) NK1 transcription factor related, locus 2 (Drosophila) snail homolog drosophila cadherin cDh6
AOUNCE CON Pregar	Ubapi under the control of the contr
# 5 5 5 F 6 7 6	microlubule-associated protein 1 light chain 3 alpha: MAP1 light chain 3-like protein 1; microtubule-associated proteins 1/1/18 light chain 3 (100% Homo saptens). neuroblastoma, suppression of tumorigenicity 1 relinoid x receptor interacting protein rip 14-1 no.5 alpha isoform histocompability 2, O region alpha hous polymerase (RNA) It (DNA directed) potypeptide 1 monocyte chemoatiractant protein 5 precursor mcp-5 small inducible cytokine member of the c-c chemokine family; strain slif a12 scya12

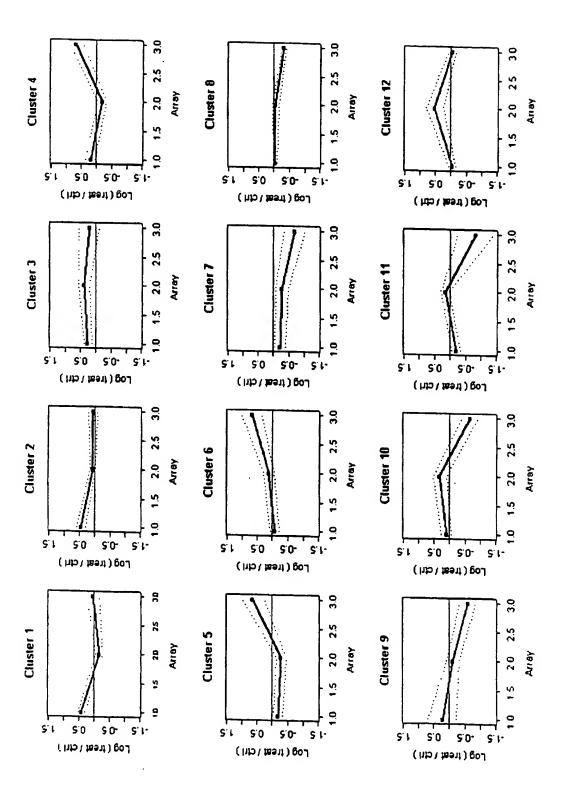
Description			aduli male lestis riken coma ckone:4930533b18			RIKEN CONA 8430432M10 gene			eloposide induced 2.4 mRNA	E74-like lactor 3				chromodomain protein, Y chromosome-like		metanocyte prolifesting gene 1		contactin 2	Jak-binding protein 1				KLKC_HUMAN Kalikrein 12 precursor (Kalikrein-ike protein 3) (KLK-L5) 70 %			sodium catcium exchanger ncx transmembrane protein	limb-bad and bear	formy benities receipt 1				TNFAIP3 inleracting protein 2		brain cdna clone mwcb-3966 unnamed protein product	COP9 (constitutive photomorphogenic) homolog, subunit 3 (Arabidopsis thaliana)	10 11 days embryo riken cdna cione:2810408b13	arginase 1, ther	dictral love homosphor			adult male liver riken cana clone:1300017a15	relinoblastoma binding protein clone mgc:6013	Appointment program McC. V.204   1/1 & Horno Saprens	to sping your entities in assuming the sping spi	COSCII DELLA FONTA MADI ICE FINA (redocina-5) maindraneferse 1A (Formita) IRNA molhidraneferse Menilla) (FINA UTasa 10 %	מייטים בייטים (קונסאויקים להודינויון וווימים אונים מספיס לא (בייווינים) (בייטים ווימים אונים אונ	cadherin 3	germ cell-specific ankyrin, SAM and basic leucthe zipper domain containing protein	nectio-tke 1	gamma-giulamyl franspeptidase ggtp	
90		Mons				71911	o O		E:24	EIC	Pcsk4			Ŝ		Myn.	?	Cutu 5			Sic30a6	Caspy			Orc2	Sic8a1	5	For	Homer	Timm44		Trip2			Cops3		Argi	ž	5		Haxa	Kopo/	91.00	200	2		Cg Gg	Gass	Nect	gg G	
Locus			75168		02611	66812	72075		13663	13710	18551		73094	12593		60315		21367	27374	72481	210148	12366	1060		18393	20541	77889	14293	26558	21856		231130		230991	26572	76371	11846	1300			15373		98/7/	2992	70802	72351	12560	74068	94332	14598	
Chai Access	2 NM 021782	Z NM 010815		2 NM 023320	2 NM 01/499	2 BC004779	2 NM 031373	2 NM 020579	3 NM_007915 3 AK003359	3 NM 007921	3 D01093	3 AK008994	3 BC017596	3 NM 009881	3 AK015017	3 AK017955	3 AK016419	3 X81365	3 AF 167573	3 AK011866	3 BC005753	3 AWAGG147	3 NM 028562	3 8000333	3 NM 008765	3 U/0033	3 AF317517	3 NM 013521	3 8C005773	3 069898	3 NM_053186		3 NM 008300		3 NM_011991		3 NM_007482	5		3 NM 033374		_	3 ARO 13243			AK008237					

3 NM 021022 3 AK011640 3 NM 009369 1 NM 009369 1 NM 011689 NM 011689 NM 011686 NM 011676 AK002682 AK002132 AK002132	110842 21413 21810 19724 75398 24017 50505 68147 24001		
AF06008 NM_016877 NM_016877 NM_013922 AK005047 AK016000 NM_011902 NM_013885 RCD40116	26440 53621 30944 52575 74457 24084 20944 29876	Psma1 Cnot4 Zfp354c Tekt2 Svs5 Clic4	20s prateasome suburil c2 pstra 1 elphe-type CCR4NOT transcription complex, subunil 4 zinc fingler protein 354C Pypothetical protein 354C Pypothetical protein 154C Pypothetical protein FLJ20432 [Homo sepiens] Pypothetical Pyp
26 2 2 3 3 2 3 3 3 3 3 3 3 3 3 3 3 3 3 3	57740 26431 77047 66229 16938 20544 83490 20465 72026	Pke Gil2 Sic9a 1 Sim2 Sim2 H47	PKE protein kinase, hypothetical serinethreonine protein Minase [Mus musculus] 100 % GITZ, MOUSE ARE GTPase-ackeriding protein GIT2 (G protein-couped receptor kinase-interactor 2) (fyro 100 % GITZ, MOUSE ARE GTPase-ackeriding protein GIT2 (G protein-couped receptor kinase-interactor 2) (fyro 100 % Similar to cytopisamic dynain heavy chain 12 954 % RLT_HUMAN 60S ribosomal protein L7 51 % Island transforming growth lactor beta binding protein 3 Island transforming growth lactor beta binding protein 3 Sobile carrier amily sodium/hydrogen rechaper member stc9a1; done image:3500839 to cell phosphoinositide 3-kinase adaptor bcap single-minded 2 Similar to TRANL_HUMAN IRNA (5-methylaminomethyt-2-thlouridyste)-methyliransterase human 89 % histocompatibility 47 RIXEN cDNA 4833424X13 gene
5 AK01938 5 5 AK01928 6 5 AK010528 6 5 AK010528 7 5 AK011528 7 5 AK01553 6 6 NM 025139 5 6 NM 02138 5 6 NM 01385 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	26401 1 1 1 1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	Melapit I Melapit I Melapit I Cidnio	mitogen activated protein kinase kinase kinase that assembly factor 1 Similar to MCM410 minicitriomosome maintenance deficient 10 (S. cerevisiae) [81% Homo sapiens] methionine aminopeptidase-like 1 A49364 Sg protein, brain - human (tragment) 70 % hypothetical protein FLJ14084 (97 % human) Similar to hypothetical protein FLJ14084 (97 % human) Similar to hypothetical protein FLJ14089 (97 % human) Similar to hypothetical protein FLJ14089 (97 % human) Similar to hypothetical protein FLJ140094 (97 % human) RIKEN CDNA 170004 (102 gene ret.HP_1600054.1 - chromosome 8 open reading frame 7 [Homo saplens] 92.94 % Interfection 12 203 subunit herre oxygenase decycling clone mgc:5802; 2a ho-2a RIKEN CDNA 430004.02 gene RIKEN CDNA 2310004.02 gene RIKEN CDNA 281009407 gene RIKEN CDNA 281009407 gene RIKEN CDNA 281009407 gene

Oene Description	Gpx\$ atulalhione peroxidase 5	_	2p3 zona peflucida giycoprotein 3		Mki67 130249 cett proliferation antigen Ki-67 - mouse 100 %	Pdg phosphogauconate dehydrogenase		flga? integrin alpha 7	Punc putative neuronal cell adhesion molecule punc	Abp amiloride binding protein 1 famine oxidase, concer-containing	edoctrome confide submit VIII 2 (59% trump)			Will modellicitisman and addition to	-	Bazi bdis-associaled zinc linger protein bazi	RIKEN CDNA 5630401024 gene	Smitar to tipase, hormone sensitive [65% Rattus norvegicus]		Nzph2 neurexophilin 2		RIKEN cDNA 2010004B12 gene			-			_	Arg arginase type ii		iks. esiitygerriesponsive iinger patiem, dane mgc. oooo	Nath Nortie disease homology				Mpv17 mpv17								.co nippostasin prostate type	b2. stosse na $*$ k* $t$ ransomino beta 2 polymentide			_			leukemia/ymphoma related factor trt transcriptional repressor ptzf		ס לאוויו ופכלקיוויותו בסוקייב אנותנייו מוכס						4b zinc finger protein 354B
Locus	14780	97 107	22788			_					78174			4 0P228			71449			18232 No		72041					_		A	22000 125		17986 Nrt			_					>		2413 HOXDS		36 1455.0	32 Ato 1b2			18 Foxn1	-		59 LA		5				Z Mikn3		4 Z(p354b
				208									22																							22 17527	!		-,	•				20030	11932			_	••		16969	50753			_		•••		27274
Clust Access 6 AK018169	6 NM 010343	6 NM 025971		6 NM 026208	6 x82786	6 AK002894	6 AK004882	86E800 MN 9	800 NN 9	6 AKD054.	6 AKD19665	6 \$45012	6 AK013903	6.563759	C 1111 007570	NA O	6 AKU1/4/U	6 AK016633	6 NM 008	6 US6650	6 NM_009133	6 AK00808	6 AK004676	600 MN 9	5 NM 013	O PASSON	6 AK00330	6 AND 1201	COVEDO MAN O	6 Degons	6 AF302138	6 NM 010883	6 NM_023371	6 BC005711	6 NM_026160	6 NM 008622	6 AK014242	6 NM 013867	6 NM 008234	O ANDIANS	21 2000 200 0	6 PCO 175E	6 600 12233	6 BC004768	6 NM 013415	6 AK002480	6 AK003747	6 NM 008238	6 AF210429	6 AK003900	6 NM_010731	6 BC004/52	6 AK016647	6 NM 009367	6 NM_007874	6 AJ006215		6 AK006923	6 NM 013744

Gene Description RIKEN CDNA 483143000   [Mus musculus] HZAM_HUMAN Histone HZA.m (HZA/m) 39 %		Acox 3 prick now assess a sexual	KINEN COUNTAGE GETTE Ofor pictrapin MAS myscalus 100 %		Cyp17a1 cytochrome P450, family 17, subfamily a, potypeptide 1	Mus musculus mmNN for hypothetical protein (ORF I) done Ricos I handed dismost a nectual protein in home Cools				Msi31 male-specific lethal-3 borndog 1 (Orosophia)	wos mostus adulti mate medicala dioritigata curva, Kitsen tila-engin ennoned library, clone:63:304 i 1024 product:hypothetical protein Kitsi kinesin family membre e3 chme imane-34:80891- kitsa.		_	Sparc - It days embryon beats frieth conda chore.School III I; Cysteine-nich grycoprotein sparc aa 1-302 Titinsto huvodhaffeal condain Mad nicht. Summan 12 46, Steine-nich grycoprotein sparc aa 1-302	LV86 hymphocyle anticen 86			Kns2 kinesin 2	Month Transition I UCZZ6442		hypothetical protein MGC13182 (Homo saperas) 81 %		Gabrina-aminobulync acid receptor mot subumi tonoropic gaba	Nmt N-myristodtransferase i		Eng edg endoglin (gr-beta receptor iii homotog: eng			MO21_HUMAN MO25-fite protein 96 %		Obc. Uniquin C				Kab33a rab33a member ol ras oncogene (amily	RIKEN cDNA 2610207116 gene		mar 17-1 offisiory receptor mar 17-1	RIKEN COLOUN'S SERVICE AND A 1700/2015/SERVICE AND A 1700/2015/SERVICE AND A 1700/2015 SERVICE AND A 1	aouir mala testis fiken cona cione; i Yuuu tegus	Gmeb 1 ghrocodricaid modulatory element binding protein 1 (Mus musculus) 100 %	MSn modes modes of the comment of th	zinc finger protein 143
Locus 98363 76383	20451		57329	13799	13074 C	12057					16582			26007				16593 #			8		30 80 82						80			98				6			e e	2			, <del></del>
							:						_		ı,									7 18107		2 13805		_	9 69008		200	66786			1933/		17254	209102	67690			17698	20841
COST ACCESS  8 NM_02889  8 NM_01178  8 NM_011778  9 NM_029596  9 AK012851  9 NM_019794  8 NM_019794  8 NM_07857	8 NM_009182	8 NM_030;	8 NM 0205	9 NM 0101	9 NM 007B	9 AJZ77212 9 NM 009751	9 AK006964	9 AK017282	9 NM_007933	9 NM_010832		9 NM_026312	9 M04429	9 AK020831	9 NM 010745	9 AKO16972	9 NM 009122	9 NM_00845		9 NM 026063	9 BC018387	9 AK020727	9 NM 023850	9 NM 008707	9 AK009578	9 NM 007932	9 NM 030718	9 NM 02767	9 NM_026908	9 NM 019519	9 NM 021875	9 AK017036	9 AK020699	9 NM 016668	9 AK017641	9 NM 024255	9 AK020444	9 AF 133300	9 AK006180	9 NM 009171	9 NM_020273	9 NM_010833	9 NM_009281

Deservation adult male lestis riken cdna done:1700082m22 dysfertin PHD zinc finger protein XAP135, isoform a [Homo sopiens] 96 % 6 days neonate head riken cdna done:5430427021 RIKEN cDNA 4920544104 gene RIKEN cDNA 4920544104 gene		consportiewhus and adenovius receptor  FAFE RNA polymerase II, TATA box binding protein (TBP) associated factor  sammal vesticle antigen. Net 2. SLP in the mammary grand; SVA-like protein in the mammary gland [Mus musculus 100 % Mist and SOK1 -related kingse  experiment and SOK1 -related kingse  cyclin D-type binding-protein 1  Li cell adhesion molecule  fRNIH MOUSE Fertian beavy drain (Ferritin H subunit) 41 %  bypothetical protein FLJ 14 11 7 (Homo sapiens) 71 %  bypothetical protein FLJ 14 11 7 (Homo sapiens) 80 %  cell line NKI 4 deMord transforming oncopere  FXYD domain-containing from transport regulator 4  creatine knase, mitochodridal 1, ubiquitous  FXYD domain-containing protein 8  RIKEN CDNA 1700039D13 gene  KLSO - LUMAN Kalikvein 8 precursor (Kalikrein-like protein 3) (KLK-L3) 78 %  cylochrome p450 cyp2d22	rein/Pi 18 13 1.1 in problekted problem FLJ 1803   Homo sapiens   71.76 %  protein cupted receptor family c group member h ginct h  ubiquilin protein ligase E3 component n-recognin 1  neurogenic differentiation 3 neurod3  complement component 4 binding protein  undifferentiated embryonic cell transcription factor utf1  plexx B2  nemediate early response 5  expressed sequence AMY1224.0  strong-leaf derived factor 4  expressed sequence AMY1224.0  strong-leaf derived factor 4  recombination activating gene 2  leucne-rich and dealth domain containing: p53 protein induced, with death domain [Mus musculus] 40.00 %  leucne-rich and dealth domain containing in p53 protein induced, with a musculus    leucne-rich and dealth domain containing in p63 protein induced, with death domain (Mus musculus) 40.00 %  leucne-rich and dealth domain mercal han mercal han mercal han mercal early mercal containing post-at mortal and polymerase if 4 14 kda subunit po2-4
Dysi Dysi Sep-07		Crade Talis Svalz Misik Misik Condbp1 Condbp1 Condbp1 Fixid Ckm11 Fixids Chm11 Fixids Cyp2422 Cyp2422	Gprcth Ubri Neurod3 C4bp Uiff Phynb2 Herpud1 Ier5 Sdl4 Rag2 Kag2 Lama2 Rpo2-4
1 73534 26903 7 72057 7 1421 70873 78807 235072			/ 4283 22222 18014 12269 22286 64209 101333 20318 101333 20318 16773 20022
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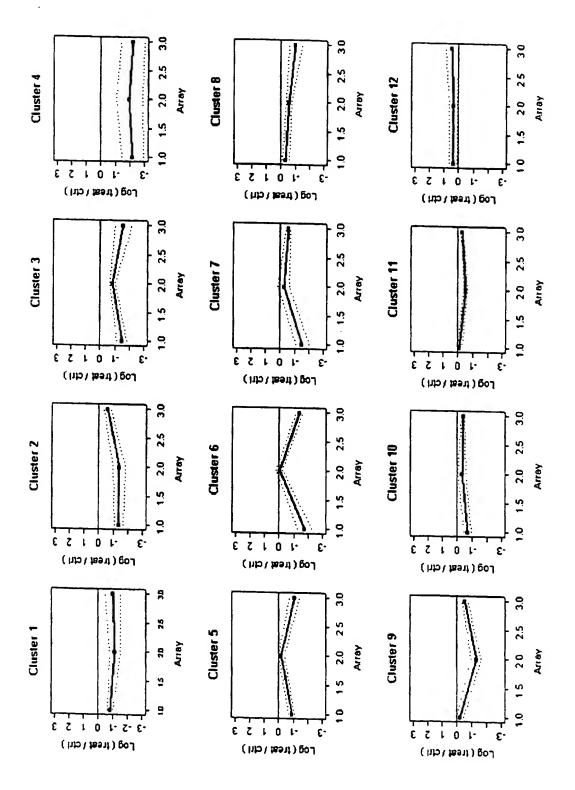


	SINGLE ASSOCIATED FORCES			And the state of t		_	_			phosphodiesterase 1C	A55575 ankyrin 3. long splice form - (28.61% human)	Ī	Ī		RIKEN cDNA C330006D17 gene	beta-amyloid binding protein bbb integral membrane glycoprotein	Wnt inhibitory (actor 1	Tetraspan NET-6 (95% human)	RIKEN cDNA 111006BE08 gene	RIKEN cDNA 2010001E11 gene	Similar to APXL_HUMAN Apical-like protein (APXL protein) huma 28 %	odd Ozifer-m homolog 2 (Drosophila)	Cabulty Campodistriction and the product Kindse	THE VIEW OF THE PROPERTY OF TH	UNA SEGNETICAT & LEVA TO USE 633, expressed	EXPLOSED SOCIAL TRANSPORT		_	_	Cado HUMAN Profein CG-100 precursor (89% human)	_	_	_	_	_		pulative seven iransementana spanning receptor puma-g pulative seven iransemental spanning receptor puma-gramman gramman gramm	Saniat to Day 100-222 - 100-222 rightenical profession Newtones - musian 62 %	Retain complex 1, acoud, gene 2	POLICIOS ACTURACION PLOTECTION CONTRACTOR CO	your most experienced as a companied from DixEN full lands benighed library where 647041701 and without benefit	RIGHT CONTROL OF STATES AND STATE	achaele-soute complex homotop-like 2 (Drosophia)	micin 10. submandibular dand salivary mucin (Mus musculus) 30 %	PGP1 HUMAN Probable pyrrolidone-carboxytate peptidase (5-oxografidase) (Pyroglutamy-peptidas 95 %		NgfA binding protein 1	caspase 9	hepatocyte growth factor	Nit protein 2 [89% Homo sapiens]	unc-84 homolog A (C. elegans)	pernala skeleja myosin kelay ovan Jeno	Cytocontic Laympian Workfeeds Special for the management of the control of the co	ACTIVITIES TO CONTROL INGOID TO INCLUDE THE CANONIC STREET OF THE	Similar to twoothetical protein FL11259 Homo sapiens! 93 %	cateium and integrin binding 1 (calmyrin)	JC6547 high suffur protein B2E - ral 37 %	atp synthase h+ transporting mitochondrial (110 comptex subumit e atp5k; lfm-1 (110-atpase
Oene	ruap).	460			Trimit	200		2fp120	ပ္တိ	Pde1c		Cdo	Mtcp1			Вър	Will					2200	3			Conf	Skrysto	Map2k6				Csnk1e	Pripm	Man2b2	8 8 8	e de la	SEE SE		7.1.5	9 5	2		Aset2				Nab1	Casp9	Ηđ		Unc843	Myze				3		Atp5k
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Chustiar Access	COCCO MAN	2000 MM 1	S AKODA154	- AKO17062	1 NM 053168	1 NM 013652	1 AK015192	1 NM_023266	I NM_010351	1 NM_011054	1 AK016847	1 NM_021339	1 NM_010839	1 AK017105	1 AK021182	1 AF353993	1 NM_011915	1 AK012571	1 AK010471	1 AK008003	AK004934	1 NM_011856	2 NW 003806	2 00017570	2 AK01/529	2 NM 011783	MM 2 17394	2 X97052	2 AK016707	2 AK014490	2 AK017114	2 NM_013767	2 NM_019991	2 NM_008550	2 NM 026031	2 NM_009757	3 NM 030/01	3 AND 3093	3 NM 010663	MM C 2406	2 AKO10946	3 AK017769	3 NM 008554	3 AK007868	3 NM 023217	3 AK007013	3 NM_008667	2 NM_015733	3 X72307	3 NM 023175	3 AF343752	3 M12289	3 NM 003843	AKONANA AKONANA	3 AK004552	4 NM 011870	4 NM_027170	4 NM_007507

Locus Gene Description 18001 igit in sinstement all actor i ecceptor igi-1 66163 Marval mischemortus inhocemos mendant la	i i		VOCCO RINER CDNA 17000061/4 gene RINER CDNA 17000061/4 gene		Get	1/44.5 ALIKE CDAA 4932403H05 gene	_	Stc35a2	•••		Kcnab3	20603 Sms spermine synthase		chromosome 20 open reading frame 10 (55% human)	Psma3	Latha	-	7 1099 Serind-Threatment funse R5581 Horne 83ptens) 44 %	Cara	Usp5	Bstt	74.239 Rabbe-linearing profesion in Management 3.38 %	NAPit	Ą	12425 Cckar cholecyslokinin A receptor	:	4 - 34 D Fulls - Fulls File States & Comment of the State of the States of the Stat	occo. Similar un injudicional programme del	Gpr 108	9	Cctea	Seroine2	Glp1r			4 III UZB A MIGROROWINT TO RECEDENCY STATES TO A CITETION OF A CITETION	710106	Bell	H2-M1	Pdcd10	3	Binlat	6 Olit 60 Olit 160 6 Olit 20 Olit 160 mitochandrial
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Description								similar to keratin associated protein 4.7 [31% Homo saptens)	Ī	9030607L17Rik RIKEN cDNA 9030807L17 oene	hymothetical protein FL 123467 (Homo camens) 93 %	tonin family 1 member 0			RIKEN CDNA 5730417B17 gene	_	c histidine rich catcium binding protein				2g complement component 1, q subcomponent, gamma polypeptide		-		ii.) UDF-GGINAC:Detaical peta-1,5-N-acety@ucosanmytransferase 3						RIKEN cDNA 2310014L17 gene	Similar to WD domain, G-beta repeat-containing protein [Homo sapiens] 83 %	RIKEN cDNA 4930528F23 gene	_	_					_							-	phosphalidylinositol glycan, dass N	2 glucose-6-phosphate deshydrogenase					-d ribaronal aratain CB biance		for hinding mentain that 130			Service to Application Flagure 1 for a notice square					_	_	hypothelical protein MGC 14827 (85% human)	sp.P46096 · SYT1_MOUSE Synaplotagrain I (Sytl) (p65) 38 %	
S. S.		A	Taries	•	1	020-020	9120		χ			Torit	5	Ceist		Alx4	Hrc	Parva	i	5	֓֞֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓		Ē	,	e sgnis	6,000	2011	5	;	<u>.</u>				DoysiS	NdufsS	S100b	Doub's	7103612			Man	CABA	Sipi	MSAB		Stau1	SixbpA	Pig	G6pd2	Code	Rn(130			Dreft and	The second secon	Ę	Neg S	200			<u> </u>		Olfre7		Dullard			
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Chuster Access	7 AK015430	2 5 7 50 MM 7	1 NM 011658	3 AKONESA	* *******	7 414 048008	NM 018888	9 AK009086	6 NM_011798	6 AK018541	8 AK007540	A A 1207743		000500 WW B	8 AK017569	8 NM_007442	8 NM 010473	6 NM 020606	* NW 007549	ANA 007574	2/C/O WN 8	00000	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	- 1444 020100	691920 MW 8	# 1103000	6 05000	8 AKU13202	a AAUISIBA	8 NM 009537	B AK009344	8 AK005105	8 AK015924	8 NM 023047	e BC002163	8 NM 009115	B AKO06201	8 NM 007565	090600 MN 6	9 AKOTA IGE	0 MM 030508	COCOCO MAIN A	9 NM 016737	9 NM 013516	9 AK012535	9 MM 011490	9 NM_011505	9 NM 013784	9 NM_019468	9 NM 009466	D NM 021540	9 AK007164	9 AK007657	A MM 010024	0 AKO18475	9 AF061744	9 NM 019825	C 4K010550	0 00012530	0 Kild 000420	DAME OCCUPANT	9 AK020272	p NM_013619	9 AK002774	10 AK012063	10 NM 026613	10 AKO21056	

Qene		2 RBT1 replication protein-binding trans-activator	Rcn2	RIKEN CDNA 5430433J05 gene		Pid	Ehd3	Slc7a1			Suit suffatase 2	Pag	Pou312 POU domain, class 3, transcription factor 2	RIKEN cDNA 4931431F19 gene	Similar to Rag C protein [94% Homo sapiens]	Sgcb days embryo riken cdna done;5730521122	Scin	Pkdrej potycystic kidney disease (potycystin) and REJ (sperm receptor for egg jefty, see urchin homologi-like	Mto1 mitochondrial translation optimization 1 homolog	_	Rixank regulatory factor x-associated ankyrin-containing protein rixank; adaptor M-1 M1	_	Polim 1 PDZ and LIM domain 1 (effin)	similar to contrapsin-tike protease inhibitor related protein (CPv.23) [Rattus nonvegicus]	Poto piccolo (presynaptic cytomatrix protein)	_	Cg10671 Cg10671 like (Drosophila)	_	_	Poir 2g polymerase (RNA) II (DNA directed) polypeptide G	Mus musculus adult male lesis cDNA, RIKEN full-length enriched library, done:1700041N15 product:CHEMOKINE-LIKE FACTOR 2 VARIANT 2	_		_		_	Eaf 1 Eaf protein	RIKEN CDNA 3732407C23 gene		_	_		Mscp milochondrial solute carrier protein	٠	CGI-30 protein (Hamo sapiens) 85 % /	Mus musculus adult male hippocampus cDNA, RIKEN full-length entiched library, clone;2900024P18 product:hypothetical Profiltr/arlengen containing profein	Similar to CDA14 [92% Homo sapiens]	divalent cation tolerant profein CUTA [Homo saplens] 91 %
Locus	11520	170742	26611	71363	72381	18701	57440	11987	69578		72043	94212	18992	70980	75220	24051	20259	18766	68291	110959	19727	94213	54132		26875	10961	68680	75803	22402	67710		17827	17175	20351	107392	11982	74427	74014	76933	12512	14429	63993	67712	13400	69740	72871	67456	67675
Cluster Access	10 NM_007408	10 AF317202	10 NM_011992	10 AK017387	10 MM_028218	10 NM 008838	10 NM 020578	10 NM 007513	10 AK009387	10 AK020739	10 AK008108	10 NM_053182	10 NM 008899	10 AK016497	10 AK015982	10 AK017781	10 NM_009132	10 NM 011105	10 AF369902	10 NM_033080	10 NM 011266	10 NM_053183	11 NM_016861	11 NM_028740	11 Y19185	11 AK014993	11 AK003939	11 AK015565	11 NM_018865	11 NM_026329	11 AK006679	11 NM_013604	11 NM_008733	11 NM 013658	11 AF233580	11 AF 156549	11 AK016628	11 AK014404	12 AK010014	12 NM_007653	12 NM_015738	12 NM_022025	12 AK019700	12 NM 032418				12 AK002828



cluster analysis t colon cancer hypothalanns

57.437 HSPC041 protein (95% Homo sapiens) 72748 hypothetical protein MGC 12904 [77% Homo sapiens]	20060 Mit2 anligen p97 (melanama associated) Identified by monoclonal antibodies 133.2 and 96.5	Msr1	i	18243 PipAa 1 protein tyrosine phosphalase type tra member clone mgc:5901; prl-1 240880 RIKEN cDNA 1200016D23 cene			14651 Glud glud gludmate dehydrogenase				Bean	15399 Hoxa2 homeebox pratein hox-1.11	19878 Rock 2 Rho-associated colled-coil forming kinase 2		19896 Rpl10a clone mgc:7602; csa-19	198 Ddit3 chap-10		25047 TAA1 manashalan sasakiin tena. 1			į	396 Git5 kruppet-like zinc finger protein glisz refated to gi subfamily; gi-kruppet zinc-finger nid myd transcription factor		548 Snta1 syntophin, addic 1		Asph			Myg1	Kail	25 Abrils 17 dans ambien head ilea calma dema 12224014.00: alrebindino casceda proteid abrils			14 unknown		Sgpp1	Chek2	46 Adpri2 adp-ribosylizansierase nad+ poly adp-ribose polymerase adpri2; 2 parb2 parp-1 like prolein parp-2	37 RIKEN CONA 2310010M24 gene		19 Cyp2u1 cytochrome P450, lamily 2, sublamity u, potypepilde 91 brain cdna cłone mncb-3966 unnamed prolein product		
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1 cc.1 coled-coll protein nucleor; partial laxp180 CAMP [Mus musculus] 100 %	2 zinc linger protein 2 zip2; mszt87	aduli male umany biadder riken cdna clone:353008 ik03 RikEn cDNA 4930542N07 gene	9 g protein-coupled receptor g2e g2a n1 CPI:anchread HTI shadish anniain 1		2004 3954 melantin-concentration becomes (400% like some continut)			Nightous (general jour Home sapens) O vicinitary recenting 40 office in	_	_	-		muserendar memorano-associated carcium-moependeni prosphotipase Az gamna - KIKEN conA 120006019 [Mus musculus] 100 %			small nuclear RNA activating complex, polypeptide 3, 50kD [64.95% Homo sapiens]		TLH29 protein precursor (Homo sapiens) 63 %		DIKEN AND STORY OF THE STORY		process types to produce assessment to the second of the s	_		mitochondrial noosontal protein L42 (Mamo sapens) 76 %	secretocranin II	refinoblastoma inhibiling gene 1	10 day old male pancreas riken cdna clone:1810061113; c-type lectin-like receptor 2 clec2	19 A M 124RA	RIKEN CDNA 1700027N10 gene	cyclin L1	RIKEN cDNA 2310056K19 gene		Property Code	ובאר אוניוע (באכ	RIKEN cDNA 2210012G02 [Mus musculus] 100 %	neuropeplide y receptor y!	balb/c asialoglycoprotein receptor mN-1; asgr1	synderan	Grant lactor ecceptor bound for the damandant seemining a throwning materials a months of	•	CCR4·NOT transcription complex, subunit 4	ATP-binding cassette, sub-family C (CFTRMRP), member 5		
23	21p2		Goiltho			Xist		01149	Mrps28	Rol8		Sic6a4	Yeldi	Scrg1					Clcn7		Protection	Src2a3		Calna		Sca2	Rbigi	Clec?			5		,	E 25	200		Npytr	Asgri	500	Caracas	03/01/0	Cnot	Abcc5		
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	(i) hunches (center) (FI 170000 Harms emissions 178 16 th	Sent	i.	5 Arn ademin	H10	Mecp2			2	Š	Rps 15			The included in the state of th	2	Cyclore C. Type in Product of Cocking and Cocking and Cocking in Security RIKEN CONA 4930566F09 (Mus mascular 55 %)		KIAA0298 hypothetical protein (human)-ribosomal protein L27a-suppression of tumorigenicty 5	5	SVs segious sequence recognistic control of the segion of	Axin exin	0.7		Alp1b3 nak-aipase beta-3 subunit alp1b3	Mich midnolin	Coned			Nos 1 nitric oxide synthese 1, neuronal		112458 hundihalical protein DKF2n5840123 1. human 97 st							Cryha2	705	EIZ4	ָבָּקָ בַּיּ	globs instituting growth and grow		rights published the fact allows any year of the control produced private and control for the control produced private and control for the control produced by the control for the control produced by the control for the control produced by the control for	Source Control and y (Source Common Galler) is sourced to receive a relation to representative specific profess (62% human)	_	Tmp2 i Iransmembrane trafficking protein	Crobit elecnitrenulativo alement bindino nodein creho t. drone impare 1500844	•	T50621 hypothetical protein DKFZp7620076.1 - human (fragment/96% human)	RIKEN CDNA 1700 129C05 [Mus musculus] 41 %	Bazi boti8-associaled zinc finger protein bazi		
74626	21654	67896		11876			2969		21892			81008		16103	77107	72747		117224	2762	268980	12005		0 C	11933	23090	74020		15212	18125		68942							1295B	724860	13663	6667	7002	22060	20067	78799	22063	68591	707.02		72519	69347	12029	74472	
7 AK015245	7 BC004630	7 AK011256	7 AK011405	7 NM 009711	7 NM_008197	7 NM_010789	7 AK010201	7 NM 025749	B NM 009390	8 NM 008740	8 NM_009091	8 NM_030733	8 AK003895 8 AB019933			8 AK013267	8 AK002385	8 AJ307670			8 AF009011	8 NM 019547	6 NM 018805	8 NM 024201	8 NM 021565	8 AK014395	8 NM_010561	8 NM 010422	8 NM 008712	8 NM 019404	8 AK004506	9 AK014319	9 NM 008508	9 NM_023587	9 NM 025788	9 NM_020579	9 NM_011665	9 NM 021541	8 ABU33513	216/00 MN 6	DAN OUBLE	O AKOOJO O		9 NM 021291			9 AK003676	9 AK017509		9 AKD12054	9 AK005/86	9 NM 007528	9 AK017033	9 BC006063

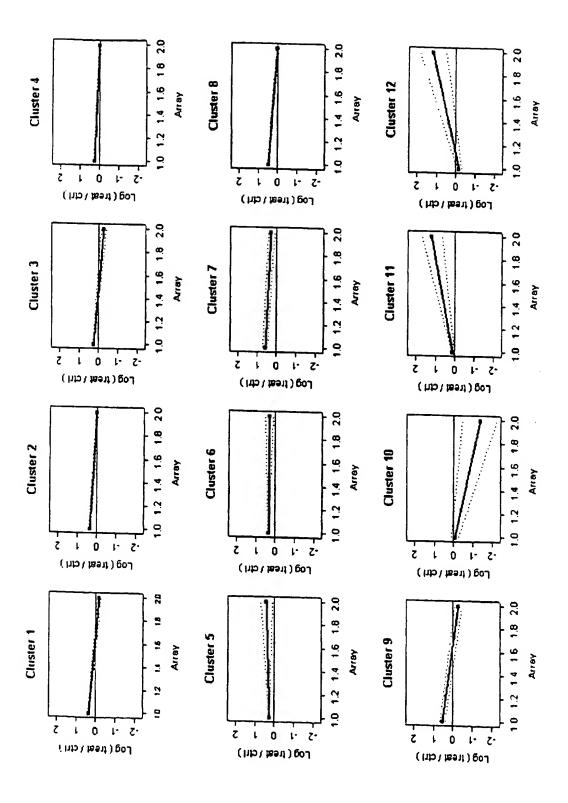
FIGURE 18.5

CGI-130 medein (2.1% Home carlene)	Cabral camp cultured of the cata-carba construction		Dahlo DARIO member 500 member 100 militari Dari da 100 militari Dari da 100 militari da 100 mi		found the ord of remainmental policies in an erane news to have a female					Pour male liver riven cana clane 130001/a15				MLNSt protein					BTB (PO2) domain containing 1 [Homo sapiens] 26.88 %	RIKEN cDNA 4931429111 ache										12 engrated 2			(1 vaccinia related kinase 1	similar to hypothetical protein FLJ10008 [Homo saptens								CODE I INDICIALITI COBI PROCETT COLOR	TO THE TOTAL POLICY OF THE PROPERTY OF THE PRO				_			hypothetical protein, MGC:7035; hypothetical protein MGC7035 [Mus musculus] 100 %	_								RIKEN CONA 1110030J09 (Mus muscutus) 100 %		_	Similar to HSPCO34 protein [75% Homo sapiens]	RIKEN cDNA 8030491N06 gene	Ī	plectin 1 intermediate filament bloding protein, 500kD (Homo segiens) 25,74 %		
											D3Ocla1			MIn51			26153182	Cachoti			Has.3		Tondeft	Tiple	Mont	Down	4	AGID		EUZ		Lag3	Vrk		Gys1	Cacna2d3	Cvn2b19	Fyved	Christ	fann	1000	2002		ć	8	Ē	Tclap2b	Fkbp	Gpr90		Cdho	}	Antho	200	ב נ	Kenmai	,	CODY			Fmo			MOR32-4		AI114950	
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zing linger prolein mkr5 3 end	exasion repair doss-complementing rodent repair deficiency, complementation group 1	cal eye syndrome chromosome region. candidate 8 homolog (human)	ignocopomina s, procopomin type s (Mais muscons) Filenting issuariatine periodolymetrials e lain 1	sulprims/erase-realed protein \$111.x1	inhibitor of growth family, member 1-tike - p33ING2 protein IMus muscalus! 100 %	polycystic kidney disease (polycystin) and REJ (sperm receptor for egg jefty, see urchin homotog)-like	origin recognition complex protein homolog morc2I orc2p yeast of replication suburnit swiss-prot acces		cysteme summing and decamony ase	polossium large condiciance calcuma-strated echamel, sublamily M, beta member 2 [Homo saptens] 95 % months his condiciance and latened and	major instructingaturung Lassas i que, a serta introduce et nu, e q'ingglori todus RZ-q'i.o RSQSAE moverin basaro de carises fermes. Namena 20 et	DISCLATING BY LOOK 14, SAME TO 11, 11, 11, 11, 11, 11, 11, 11, 11, 11	THE TOOL ASSOCIATE OF SECURITY	Add059 seriel escretor protein to receive a mouse than and 100 %.	neutrophi cytosolic factor 4	similar to -pir:131100 - 131100 probable potessium channet 2 - rat 89 %	inler-atpha trypsin inhibitor, heavy chain 2	ATP-binding casselle, sub-family G (WMITE), member 2	lesticular cell adhesian molecule 1	HIKEN CINA 290055.279 gene	prospriodantean (Mus musculus) 100 %	outsi vagaine entro in antipore e a Soute e antipore e e e degeneración esta entro esta entro entro entro entro esta esta entro de entro en	FINESCO. CONTROL 2310032016 gene.	midine 1	smcy selected adna an the y	RIKEN CINA 8430431K14 gene	WATER DESCRIBE SERIO FOR A RECENT OF A RECENT OF THE SERIOR OF THE SERIO	ווייין בין יוין ליבין יויל בין ליבין יויין בין ליבין און בין בין ליבין און בין ליבין און בין ליבין און בין ליבין	metabolimate recentor lorse 1 (Glu 1)	RIKEN CDIA 11 10025612 gene	phosphotipase C, zeta 1		neurolysin; neuratensin endopeptidase (90,33% Homo sapiens)	similar to RIKEN CONA 1810008A16 gene [Homo sapiens] 64 %	platetet grycoproken uparpita membrana raceptor suburat von watebrand lactor receptor surface	nuclear matrix cretein n84 (95% human)		erbb2-interacting protein erbin basodaterat	RIKEN cDNA 4930473B18 gene			pollovirus recentor related 3 -vectin-3 biblia: nectin-3 camma (Mus musculus) 100 %	mast-cellige high affinity receptor to-epsilon-ri alpha subunit precursor			Similar to Voltoge-dependent calcium channet gamma-4 subunit (Neuronal vottago-gated calcium channet) (28% Human)	centramere autoantigen C	mus inostable aludi mate feats a Contra, Anta-tra marengui emis en iona appi, come i contrasementi entre ana so disciplis phosphia appia-naeelyducosammephosphotamilerase 2 dpagi2	calhepsin 3
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phorbol-12-myristate-13-acetate-induced protein 1; Noxa protein [Mus muscutus] 100 % bone morphogenetic protein 1	pulative N-acetyltransferase Camello 4; RIKEN cDNA 0610037016 gene [Mus musculus] 100 %	secreted frizzled-related sequence protein 5	RIKEN cDNA 4930556/24 gene	nuclear pore membrane phycoprolein 210 pom 210	zinc finger protein 354C	RIKEN cDNA 4930406624 gene	RIKEN cDNA 4632412124 gene	lymphotoxin A	C-type (calcium dependent, carbotydrate recognition domain) lectin, superfamily member 6	msx3 the drosophila melanogaster muscle segment homeobox msh protein encoded by genbank accession number u33319
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Description RS4 HUMAN 40S ribosomal protein S4, X isoform (Single copy abundan) mRNA protein (FCR10) 93 %	e of (1906) (respectively) and the original state of the original state original state of the original state o	oduli mele smali intestine riken cdna clone:201032016; mpi3 milochondrial ribosomat protein i3 i3mi	teriboa bendred 24 mRNA	cori elidoco o Romano activa				fritziad homolog drosophija fzd 10	translocase of inner mitochondrial membrane 13 homolog a (veast)	immunoqtobulin superfamily member igst6	pescadillo homolog containing brot domain zebrafah pes 1		lymphotactin		adult male liver riken cdna clone:1300017a15			aspartate-beta-hydroxytase asph; aspartyl beta-hydroxytase	Ravin containing monooxygenase 1	sutotransferase-related protein SULT-X2		estrogen receptor bela estra erbeta; esr2	The fines retibiling R	בייה וויונים היינים או				nucleobindin 2		17 days embryo head riken chae clone:3200001111; cysteine-rich phycoprotein sparc as 1-302	histone deacelylase 5 hdac5 class ii	lipotransin hormone-sensitive lipase hst-interacting protein						receptor tyrosine kinase-tika orphan rort - ROR1_MOUSE Tyrosine-protein kinase transmembrane receptor ROR1 precursor (Neurotrophic tyrosine kina 100 %			RIKEN CDNA 2410012H22 gene	BMX non-receptor tyrosine kinase	homeobax pratein hax-1, 11			- should be a city and	g equal y southing pan innelin membrane channel eretirin altha 10		obsendinase A2 orono IIO				150621 hypothetical protein DKFZp7620006.1 - human (fragment/R96% human)	ggele lor odorani receptor Artistica. Bele lor odorani receptor Artistica in del primi procedite inhibita proceda Maria marenia del Basta mismo del primi del proceda del primi proceda del prim	מבנים. ומים מסברו שנילו מיביר. ממופט חו שימיו. לע מספר יו שומיו. לע מספר יו שימיו. לע מספר יו שימיים לע מיביר יו	ribosomal protein s15 rps15			dentin statophosphoprotein
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Description achit main unnary bladder riken cdna clone:95330091k03		oocyle specific homeobox 1					avaZ zwogene	Schrift (Winds China Chi	astrolactin neuronal migration protein go 14		RIKEN CDNA 2700023X09 gene	Injeatite motif protein trim26 alpha	protein lyrosine phosphatase, receptor type, O	simple reveal semisones-containing transcript (28% Mas musculas)	DATE I market before with the second of the	CODE, Included Code Code Code Code Code Code Code C	netrometan					дзагуте В			zinc finger protein of the cerebellum zict		RIKEN cDNA 4930441014 gene		• 138487 tastin • human 51,90 %		hypothetical protein, BC004044; hypothetical protein MGC7673 [Mus musculus] 100 %		iloggm upregulated with neuriegrown in neuropiasiona ceus, mesencifina siem ceu protein uscoz crane ingeli oko				interferon concensus sequence binding protein 1		A.C. 1 Synapsis is - Sea in a	Sulforansierase-realed progen SUL 1-X1	RIKEN CDNA 24 10004F06 gene	peroxisomal biogenesis factor 11a	108675 hypothetical protein DKF2p564F0522.1 - human (fragment) 49 %	10700 and a solution of the so	SKOT PROBED DUBLING THE SECURITY FOR SECESSION THE TOTAL SECURITY FOR SECURITY FOR SECURITY S	Anice en in an anice en in anice en in an anice en in ani					hundihalital nitrialin El 110560 (82 88 Homo saniens)					nt-3 neuratrophin-3	Mus musculus adult male testis cDNA. RIKEN full-length enriched fibrany, clone:4921510J17 product:hypothetical EF-hand containing protein	ISK4_MOUSE Serine pratease inhibitor Kazal-type 4 precursor (Peptide PEC-60 homolog) (MPGC60 protein 100 % /	erythrocyte protein band 4.1-like 3	Similar to 148208 zinc finger protein 30 - mouse 44%	CGI-127; yippee protein [100% Human]	cylochrome c axidase, subunii VI a, polypeptide 2	proges(grove membrane binding protein [73% Homo sapiens]	
18558		7 1468 Obox 1					2457 C2C27	70/01			72569	22670 Trim26	19277 Ptpro	104479	1070	1954 4951	DELIG NATO					14939 Gzmb			22771 Zic1		68271		78733		80752		83485 Ngrn				15900 Icsbp		76103	57429 Sult-x1	1881	18531 Pex118	73833		20479 Vps4b	Sole along					17577	200				18205 Ntf3	70894	20731 Spink4	13823 Epb4,113	74352	69687	12852 Cox6a2	70804	
Cluster Access 1 AK020653	1 NM 080433	1 AK018362	1 AK015384	AROTOBA	OCOLLO MINI	AR019/32	000500 MM	1 MM 100483	I NM UUTABS	1 AK018386	1 AK012283	1 AF230395	1 NM 011216	1 BC001203	STATE OF STATE OF	OFFEDO WAY	CICRIO MM I	AROOSOSO	1 AK007941	1 AK009330	1 AK008724	t NM 013542	1 NM_026312	1 AK005095	1 NM_009573	1 AK005200	1 NM_026649	1 NM 007753	1 AK021408	1 AK004155	1 NM 030565	1 NM 025424	NM 0313/5	1 NM 007685	1 AK011461	WW 025989	NM 008320	1 AK010336	1 AK018054	1 NM 020564	1 AK010386	1 NM_011068	1 AK003496	1 NM 017465	1 NM 009190	DO TO THE TOTAL	AKONACAB AKONACAB	* *******	1 AKOUSSS2	CESSON IN I	1 01017305	SOLUTION I	1 NM 019477	1 BC017625	1 NM 008041	1 NM_008742	1 AK014861	1 Y11505	1 NM 013813	1 AK014610	1 AK010201	1 NM_009943	1 AK014543	

Description nuclear receptor coactivator 4	RIXEN CDNA 2810489J07 gene	hypolhelical protein FLJ12691 [81% Homo sapiens]	HSPC 166 prolein [Hamo sapiens] 89 %	BRCA1 associated protein odd outline and odd outline the homelon 3 droscothia odd3	T50839 UA/U6 small nuclear ribonucleoprotein hPrp3 (imported) (99% human)	RIKEN CONA 2510027N19 [Mus musculus]		ribonuclease 6 precursor (Homo sapiens) 67 %	Gride burding protein 2 tobbs	unesm arany mentroes v HSF4 MOUSE Heat shock factor protein 4 (HSF 4) (Heat shock transcription factor 4) (HSTF 4) (mHSF4) 33 %	oxysterol binding protein-like 3	integral membrane protein 2A	secretory carrier membrane protein 5 scamp5	Safrogytean, alpixa (bytarophine) Saboraled glycoprotein)	e co face in occidentations	chromaffin granule atpase ii homotog bos taurus encoded by genbank accession number u51100	6-pyruvoyi-tetrahydropterin synthasefdimerization cofactor of hepatocyle nuclear factor 1 alpha (TCF1)	neurotrophic brosine kinasa, receptor type 3	GA repeal binding protein, Deta Z, pr. 25.3530 · A5.3530 transcription lactor GABF beta Z-1 chain · mouse 100.00 %	integrated potential to the control of the control tearing the con	בנוסי בנוסי בתרוני לאים אפנים	adenosine deaminase, IRNA-specific 1 (Mus musculus) 100 % /	inactive X specific transcripts, gene with no protein product	RIKEN CDVA 9130015A2 gene	KIKEN CDNA 4950406EZ4 BENB	hypothetical protein FLJ12547 (38% human)	refinoid x receptor interacting protein rip14-1no.6 alpha isoform				organic acts control acts contr	to be a locus from bases 250554 to 501917 section 2 of 3 the	chlaride channel pratein clcn3	RNA binding motif protein 5	1423/2 probable guary/ale kinase (EC 2.7.4.8) 1, membrane-associalica, splice form b - mouse 44 %	arcinine vasconessin receptor to avor 1b: hoe	milochondrial capsule setenoprotein mosp		hypothetical protein HSPC 194 [Homo sapiens] 85 %	myeloid/lymphoid or mixed lineage-leukemia translocation to 3 homotog (Drosophila)	ATP-binding cassalte sub-family B (MDR/TAP), member 10	zine finate poletin 46 zip 146	ghulamale receptor ionotropic ampa1 alpha gria1			рако	** ***********************************	actin Federal Diophers, Subtrill 1A	aspary annayepunase pir: 14768 - T14768 hvodhelical protein DKF2o566K1924 1 - human (fragment) 93.58 %		
Locus Gene 27057 NC084	72815	71685	46501	72399 Brap 23965 Odz3	19101	67711	67590	68195	14276 FOFZ	71066	71720 Osbpi3	16431 Ilm2a	56807 Scamp5	40391 Sgc3	20,000	11980 Atp8a1	13180 Pcbd		213054 Gabpb2		CBIO	30947 Adat1	213742 Xist	71581	/585/	77749	20186 Nr 174			06476 thurs	55925 SWR	269846	12725 Clcn3	83486 Rbm5	52357	26361 Avertb	17235 Mcsp		66154	70122 MIR3	56199 Abch10	26465 Zfn146	14799 Grin1			18511 Pax9		Se443 Arpcia	1343/ Unpep 70255	2520	
Cluster Access 1 NM 019744	1 AK013453	1 AK016622 2 AK002747	2 BC010495 2 AK007640	2 NM_028227 2 NM_011857	2 80006782	2 NM 026330	2 NM_026260	2 AK015947	2 AKONEBEA	2 AK016803	2 AK004768	2 NM_008409	2 NM_020270	2 AKMA171	2 AK020972	2 NM_009727	2 NM 025273	2 NM_008746	2 AK020406	2 NW ORRYG	2 AK01 1405	2 NM 013925	2 X59289	2 AK018624	2 AK015107	2 AK020339	2 NM 009108	2 NM_021437	2 NM_020508	S MACCO	2 NM 018802	2 AE000664	2 NM 007711	2 BC003988	2 AB045323	2 NM 011924	2 NM 008574	2 NM_011086	2 NM_025387	2 AF333960	2 NM_008621	Z NM 011980	2 NM 008165	2 AK005864	2 NM_018819	3 NM 011041	3 NW 010587	3 NM 019767	3 AKD19934	3 NM 026437	

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Description	mitogen artivated protein klnase kinase 1	aduli male tung riken cdna chone;120001715	exchandered kings 2 Muse musefuls 100 %	adult male stomach riken odna chare 22 tok 15d 9: te channel	Carda fune homes has sets	BIKEN COM 1700'N IN Cons	בייברי בייני	T cell activation increased tate extression		perisin DESTACESIONA 1. human all manuscript by human	process on a possible of the first of the fi	ChAZ CHECKDONI normong (s. pompe)	Sry-box containing / 60x/	open reading frame 61	cdc2/cdc28-like protein kinase 4 clk4 sr; cdc like clone mgc;747?	RIKEN CDNA 2010009J04 gene	RIKEN CDNA 5730455013 gene	poliovirus receptor-related 3 -nectin-3 arata: nectin-3 beta: nectin-3 camma (Mus muscrimus) 400 %			anithalial medain 2	Carburola in Licinal Process	consumptions to the contract of the contract o	receiptor involvers traces - 5 era bitatives, crime produce conceptor men	Seture (a Cysterie p Dottalese minimizer, Gode O mentage 4 hardelalisation of seture CBB (12) Harde analysis (C)	Informational Formation States of Frontier States of Administration and Administration and Administration of Administration and	South transcription to the Children Could be a controlled to the Could be a controlled to the controlled to the could be controlled to the controlled to the could be controlled to the could be controlled to the controlled to the could be controlled to the controll	opporting a constraint (Springer)		discs bross homelen 1 (Prescentia)		sarcogiycan gamma 35kd dystrophin-associated giycoprdein sgcg	monocarboxyalate transporter	insulin receptor tyrosine kinase substrate (90% human)			Mus musculus adult male testis cDNA, RTKEN full-length enriched fibrary, done:4930485412 product:hypothetical protein			organic cation/carniline transporter ocin2	adult male cacum rixen cdna clone:9130012o13	ribosomal protein 113a rp113a		3-beta-hydroxysteroid dehydrogenase/detla-5-detla-4-isometase; hydroxysteroid dehydrogenase-6 detla<5>-3-beta hsd3b6		•	WO repeat contain to	giulanat acto decarooxylase i	CAN'N LIMAAN CARISTO TO 1040 Mineral	SNAJ TOMAN SOUTH RESID TO COATE TO COAT	THE PARTY OF THE P	KINEN CUNA JULIO BETTO POLI 2 MOLICE Deliveira sakida PD1 redescribin El Bauerte Protectibles - Endonielessell 32 & Minuscribus	ב כרג" שייכיסר אבייסגוניסי בשנים בי כר ליסגונים ליסגונים אבייסים של שייכיסים אבייסגוניסים אבייסגוניסגוניסים אבייסגוניסים אבייסגוניסגוניסים אבייסגוניסים אבייסגוניסגוניסגוניסגוניסגוניסגוניסגוניסגו	prefettin 2 pidn 2	rab 3a member ras carcopene family	GPV_RAT Platetet glycoprotein V precursor (GPV) (CD42D) 30 %			cop9 constitutive photomorphogenic subunit 7b arabidopsis cops7b	cystatin 8 (cystatin-related epididymal spermatopenic)	ADP-nbosylation-like factor 6 interacting protein 5		
Locus Gene	26395 Map2kt	7 1740 Pvr14	12566 Cdk2	16509 Kcne1	12592 Cdr4	73344		84544 Tactile		67002	20000	Social Creek		216157 Orf61	12750 CR4	76523	70567	58998 Pvr13			1771 Emn?	56264 Covers			71793	74068 Coss	2072B Spir		1202 Appl	13383 Dloh1		24053 Sgcg		66839			74989			20520 Stc22a5	71597	22121 Rpl13a		15497 Hsd3b6			61896 War 10	1000 0101	70300	10020	24102	70770		18637 Pidn2	19339 Rab3a	06650				13012 Cs18	65106 Art6up5		
Chuster Access 3 NM 018767	3 NM 008927	3 AK004821	3 NM 016756	3 NM 008424	3 NM 007674	3 AK006598	3 AK017516	3 NM 032465	3 AK005050	J AKNIJUIZ	1 MM 010481	2000	S NM CI 1840	3 80005494	3 NM 007714	3 AK008163	3 AK017666	3 NM 021495	3 AB047557	3 BC006075	3 AFORTRZB	3 NM 019696	1 NM 009587	NW COOLSE	3 AKOUR74	OCCUPANT C	3 NM 011461	3 AK012248	3 AK004627	3 NM 007862	3 NM 008834	3 NM_011892	3 NM_009196		3 AK009683	3 AK004148	3 AK015502	3 BC018324	3 NM 023181	3 NM 011396	3 AK018615	3 NM_009438	3 AK016624	3 NM_013821	3 AK007368	3 MM U23933	3 AF 290073	7 0000 MM C	•		בינטאטאר ב	3 AKD14354		3 NM 011070	3 NM 009001		3 AK005756	3 NM_008516	2			3 NM 029682	3 NM_007990

Cluster Access	Locus Gane	Page rivilian
4 AK020379	77205	Mus musculus adult male diencenhalon cONA. RIKEN full-lennth enurhod threats status accessorate about accessorate
4 NM 022324	64136 Sdf211	stromal cell-derived factor 2-like 1 Mus musculus 100 %
4 AK016101	75266	S66619 Mas 20 protein - (42% human)
4 AKD08069	0/869	RIKEN cDNA 2010003119 gene
4 NM_016700		
4 NM_007888	13543 DW2	dishevelled 2 dsh homolog drasophila dv/2
4 AFT 16530	50/64 F5r615	F-box only protein 15
4 NM 009211	20588 Smarcc1	SWI/SNF related, matrix associated, actin dependent recitation of chromatin subtainty member 1
4 AK019034		
4 302995	11350 Abi1	testis-specific c-abl protein
4 M14872	14714 Garh	gnrh-gap encoding gonadotropin-releasing hormone and gn-rh-associated peptide gap precursor
4 NM_008076	14409 Gabrr2	gamma-aminobulyric acid gaba-a receptor subunit tho 2 gabrr2
4 NM 008874	18797 Picb3	phospholipase c beta3
4 AK014534	70802	DM3A_MOUSE DNA (cytosine-5)-methytransferase 2A (Dmmt3a) (DNA methytransferase MmulliA) (DNA MTase 30 %
4 BC003914	68585	3-hydroxyisobulyrate dehydrogenase clone mgc:7270
4 NM 020591		
ZBCSOD WAY		
4 AK005118	7,1860	KIKEN CONA TOOUGH DA gene
4 NM D08113	14570 Arhadia	tryponrence) potent tractor use to separate superior control to the control tractor of transmission of tractor of transmission of tractor of transmission of tractor of transmission of tractor of tracto
4 NM 010877	17970 NC/2	The year of the second
4 NM_031260		
4 NM 008644	17830 Muc10	mucin 10. submandibular gland salivary mucin
4 NM 008904	19017 Ppargc1	peroxisome proliferative activated receptor, garmma, coactivator 1
4 AKO16435	Seeby Perz	peroxin Z. peroxisomal largeling signal 1 receptor-like; KliKEN EDNA 1700016J08 gene [Mus musculus] 100 %
4 AK002581		
4 NM_016705	16564 Kit21a	k1/21a kinesin-like protein
4 NM 009674		
4 NM_013762	27367 Rpd3	) i protein yeast ribosomeil 13 homologue
4 AKONGA 10		
4 AK010364		
4 AB008183	12943 Pcdha10	protocadherin alpha 10
4 NM 020493		
4 NM 026096		
4 NM 010484	15567 Sic6a4	serotonin transporter sert neurotransmiller; encoding serotin
4 MM 008411	16433 Ilmap1	integral membrane-associated protein 1
A AKOOAS32	74 105 Gast	dokenyu-ispasopyooggostaanana kayaten gyotogranasersassa andri sennistad memos adabka asa nedalisina ADE kindina anaksis 3
4 NW 024263	2000 COLE	עניע מספרביפני, עם וויים מספףווי במו בחומויווין. חחר טוומייון אחר מו ממיים
4 AK017922	74752	Mus muscutus adult male thymus cDNA. RIKEN full-length emiched library, clone:5830412H02 product hymothetical protein
4 AK017044		
4 NM_009871		
4 AK017598		
4 AK016452		
4 NM 029082	72000	KINEN CLINA SEGUAL INC. Gene Adult media bisocuramente alton celebra-DOMINO Ant
4 NM 012053	26961 Rolls	adon ner eight. Ribosomal protein LB
4 NM 010925	18114 Nnp1	novel nuclear protein and t
4 NM 008191		
4 L25890		
4 NM OORGRO	19262 Pinra	terirocke common antinensetated observations or enumer
4 AK010169		
4 AK013041	69940	Sec3-fike; hamolog of yeast exocyst protein Sec3p [Homo sapiens] 96 %
4 AK006168		
4 NM 007516	11901 Harad	hatarmanaka audasa ahkanadanyadain D
4 NM 017471	odnu issi	וופופו מלפו ופרסים וורכופים וורכפים וורכופים וורכפים וורכפ
4 NM_010700	16835 Ldr	fow density lipoprotein receptor ldfr
4 NM_010060	13411 Dnahc11	dynein axon heavy chain 11 dnahc11
4 AKU15948	75196	RIKEN CUNA 493033ZLZO gene

Description	מים והסיימות והפים וואה ביים ביים ביים ביים ביים ביים ביים ביי	prop. 4 arydrovyskae aproblastovovní muscha specific gene i Homo sapiensi 99 %		Mus musculus RIKEN cDNA 2700063A19 gene, mRNA (cDNA clone MGC:6532 IMAGE:3492667), complete cds		Mail 1 registed chloride channel	ven nivov givov grindin displaces, related en 20. relativa bindina interfinal rei	לו כוכנון ביותר בי	prolein (peplidy-proM cis/trans isomerase) MIMA-interacting, 4 (parvufin)	very low density lipoprotein receptor	dba/zi della profeasome sutumii	Similar to hypothetical protein FLJ20211 [80% Homo sapiens]		G7e protein	adaptor-related protein complex ap-1 mu subunit done mgc:6219; clathrin-associated adaptor medium chain ta ap1m1	RIKEN cDNA B430319H21 gene	relinoblestorna binding protein 4	KIKEN CUNA 17 UNUSGCU3 gene	For the desired to the form	Productive commensor reco	RIKEN 201000HOB ome	(br2):-box-containing	12 days embryo embryonic body between diaphragm region and neck riken cona clone:3430068a13; ribosomal protein s5	macrophage scavenger receptor type I; sr-a	interferon response element binding factor 1	translocase of inner milochondrial membrane 8 homolog a yeast done mgc:67:30; timm8a	v-el simila fettemala vital orogène homolog B (13s fetaled)	hypothetical profession to LLAZ 1906 [16% March Sagletins]	defense/intributly protein activity, mudeal inter-pyllicia z.cbroxygaliciae activity		KLKC HUMAN Kallikrein 12 precursor (Kallikrein-like protein 5) (KLK-LS) 70 %	copine IV	NADH dehydrogenase (ubiquinone) 1 apha subcomplex, 3	Then code 28 100 JBK 1971k	Bipna-4-durandase raua			interrin aloha e eoithelta-associated Itgae	mannosidase alpha man la		ADP-ribosylation-tike factor 6 interacting protein		T47144 hydratheirs inclidin DKE26/6151347 1 - himan (fragment) (40% himan)	adult male bus riten con a done 12000 3009; partial roquois homeobox projein ix5	confacili associated protein-tike 2		RIKEN CDNA 1810042K04 gene	KIAA0598 gene product (92% Hamo sapiens)	Msx-interacting-zinc tinger		AD21 MOUSE ADAM 21 precursor (A disintegrin and metalloproteinase domain 21) (ADAM 31) 100 %	brain protein 16; DNA segment, Chr 15, ERATO Doi 741, expressed (Mus musculus) 100 %	nuclear receptor sublamity group a member ru5a t	strate tim homotony domain rentaining (amily R (averline) member 2	percent intringing formation containing and programme processing the more processing and processing the processing the processing and processing the processing and processing the processing and processing the processing the processing and processing the processing and processing and processing the processing and processing the processing and process	
Locus Gene	20000 1000	73830				229725 Mctc	12304 Cai	1000	69713 Pin4	22359 Vidir	19175 Psmb6	72341		110956 g7e	11767 Ap1m1	77849	19646 Rbbp4	18654	16160 14000	45 190 Dist	72152	13813 Eomes	20 103 Rps5	20288 MSr1	26573 treb/1	30059 Timm83	64 143 Rath	72651	90111 97FQ1		69511	74020 Cpne4	66091 Ndufa3	72699	15932 1609			16407 Itase	17 155 Man1a		54208 Avi6ip		21164	54352 hrs	66797 Cntnap2		90599	77590	17344 Miz 1		56622 Adam21	59053 Brp16	26423 Nr5a1	CANADA DECEMBER	17434 Mocs2	
Chuster Access	Carolina P	4 AK004664	4 AK015769	4 BC003972	4 AK015864	4 BCD03247	4 MM 009787	4 NM 021430	4 AK010338	4 NM 013703	4 NM 008946	4 AK011270	4 AK013485	4 NM_033075	4 NM_007456	4 AK021026	4 NM 009030	5 AKOTBBB1		5 SC00232	5 AK011486	S A8031037	S NM_009095	5 AF203781		S NM 013898	5 NM 022327	S NM 028311	2 NM 008382	5 AKO14759	5 AK009217	5 AK014396	S AK005243	5 NM_023684	S NM 008325	5 AK/00114	S NM DARK	5 NM 008399	5 NM 008548	5 NM 010174	5 AF223953	5 AK007715	5 AKO16865			5 BC003954	5 AK012616	S AF320786	5 AK018146	5 AK012661	5 NM 020330		S NM 008050	5 AK009120	5 NM 013826	5 AK002327

Description Chromosome 20 open reading frame 108 [Homo sapiens] 77 % DMEZPA34C254 potent himon sapiens] 90 % RIKEN CANA 49304 71.421 gene relinal degeneration, stow (retinitis pigmentosa 7) cortaction tith contaction tith eutharyotic rearestation initiation factor 2 alpha kinase 3 \$ 10151 transforming protein tith - (34% mouse (strain balb/c)) paired like homeodomain factor 1	amiloride binding protein 1 (amine oxidase, copper-containing) odorant receptor S18 gene it type c glycoprotein mcg	pregnane x receptor pregnanc-activaled nuclear vanin 3 prosphoprotein enriched in astrocytes 15 RIKEN cDNA 9530004P13 gene sy-box containing 5 sox5 NKT transcription raciated, locus 2 (Drosophila) N-myristoyfiransferase 1	RIKEN cDNA 2010309E21 gene grawh hormone reteleasing cd 1 integrin alpha v subunil vitronectin receptor CD8 antigen, beta chain matrilin 3 RIKEN cDNA 2310046K23 gene steroid sensitive gene 1 jiunctophilin 3; junctophilin type 3 [Mus musculus] 100 % Mus musculus gene for Cox17p, complete cds cytochrome P450, family 17, subfamily 9, pobypeptide 1	SRY-box contining gene 14 translocase of outer mitochondrial membrane 40 homolog (yeast) zic3 protein zic3 protein capping protein actin filament muscle z-line beta ctone mgc:5082 protein inhibitor of activated STAT 3 serva domain, immunolopholina domain (fg), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4G 112514 hypothetical protein DKFZQ43073.1 - human 74 % hypothetical protein FLJ10559 (100% human) dylamaie receptor channel subunit zeta-1	ctone or 1.72m 13 oifactory receptor orthologous to or 1.72 RAS-related C3 botulinum substrate 3 doublesers and mab-3 related transcription factor dmrt1 candidate sexual regulatory protein; franscript src homology three (SH3) and cysteine rich domain (Slac) est2 repressor lactor erf
Locus Gene 67017 70235 75810 19133 Rds 19133 Rds 19265 Eif2ek3 71246 19127 Prop1	76507 Abp1 56860 Ors 18 56315 Rhcg 83434 Rsh11	26464 Vnn3 26464 Vnn3 18611 Pea15 77363 20628 Son5 20231 Nkr1-2 18107 Mmt1	66488 1440 ligav 12526 Cd8b 17182 Main3 76968 67895 Ssg1 57340 Jph3	22733 Torm40 22773 Zic3 22773 Zic3 12345 Cappb 1 229615 Pas3 26456 Semo4g 77718 77718 15356 Hrngcl	Unknown 170458 50796 Dmrt1 20840 Slac 13875 Ert
Chulw Access 5 BC016210 5 AK011054 5 AK0115544 5 NM_008938 5 NM_001021 5 AK017101 5 AK017101 5 AK017001	5 AK005423 5 NM, 020290 5 NM, 020596 5 AK020727 5 NM, 028596 5 NM, 031255 5 AK005168 5 AK005168	5 AK012814 5 AK01887 5 AK01987 5 AK009912 5 AK005205 5 NM, 011979 5 NM, 008556 5 AK02054 5 NM, 011444 5 NM, 009173	5 NM_052686 5 NM_052591 5 NM_010285 5 NM_010285 5 NM_010286 5 NM_010770 5 NM_011349 5 AK01985 5 AK01985 5 AR01728 5 AR01728 5 NM_050505 5 AR01728 5 AR01728		6 AK003882 6 NM_020500 6 AB040819 6 BC004674 6 NM_016853 6 AF359382 6 NM_01055

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Describtion riken of na 1110004107 1110004107/ik RIKEN cDNA 2310068104 gene	146537 transcription factor 1, neural - rail 41 % esophageal cancer associated protein [Horno sapterns] 92 %	RIKEN cDNA 4930451E06 gene 10 11 days embryo riken cdna clone:2810408b13	Similar to 148668 zinc finger protein 51 - mouse 43%	prolein Z. wiamin K-derpendent plasma glycoprotein	lar encoding protein tyrosine phosphalase; ptarf	amin cacyfase 1	paniothenate kinase 1 lymphocyte antigen 78 ly78	SREB3	secretary protein by8	RIKEN CDNA 4933428119 gene	KINEN CINA 5) 344000304 gene cyclin-dependent kinase inhibitor 2B (p15, inhibitos CDK4)	Mus muscurus adur mare testis cumA, KiKen full-length enriched library, clone:1700058G18 product.hypothelical protein cerebellum zic4 protein	M.musculus ORF1 and ORF2 genes	phosphotipase Chike 2 odorant receptor St gene (Mus musculus) 100 %	nuclear receptor subfamily 0, group B, member 1	FAFT KNA polymerase II, TATA box binding protein (1847-essociated factor		ggamma-amindbulktyr, caki (1444-A) feceblor, subunit bela 1 hes-3 helis-leon-helis transcrintien factor	18 days embryo ilken cdna clone: 11 10008f24; alpha-actinin		proly endopepiidase prep	DAL: mDNA binding section	The state of the s		prolein phosphalase 1, calalytic suburit, garmua isoform	RIKEN CDNA 3110079015 gene	Cadherin	solute carrier (amily 39 (zinc transporter), member 3		inclinger protein 162	Straight in Intersection 2 176% Home senions	docamine receiptor 2	RIKEN CDNA 4933424G06 gene	smilar to hypothetical protein [Mus muscadus]	protocoaterin apna 10 - reivir-usi raz. 1 - protocaanem apna 4; caanem-telatea neuronalreceptor 1 (Mus musiculus) 81 9 FKSG6 binding protein 11
Locus Gene 66142 Cox7b 69653	71248 71517	74858 7637 I	67607	2014 F0809	19268 Mpd	109652 Acy1	75735 Pank1 17079 Ly78	70771 Sreb3	50501 Prok2	71198	12579 Cdkn2b	22774 Zic4	000000	224850 PICIZ 56858 Ors1	11614 NrOb1	240/4 181/		15207 Hes3	11472 Acin2		19072 Prep	66870 Pairhai			19047 Poptcc	73234	t2565 Cdh9	106947 Slc39a3	;	22668 Zip162	74855	13489 Drd2	71166	0000 Decker 00	66120 Fkbp11
Cluster Access 6 NM 025379 6 RC/05635	6 AK016974 6 AK011615	6 NM_016855 6 AK015439 6 NM_029583	6 NM_033560 6 AK01.1166 6 NM_02624	6 AK008575	6 NM 019746 6 NM 011716	6 D29987 6 NM_007841 6 NM_025371 6 AK017419	6 NM_018798 6 AK019493 6 NM_08533	6 AK003099 6 AK014326	6 AK019640	6 AK016972	6 NM_007670	6 NM_009576	6 X98456	6 NM 020288	6 NM 007430	6 AK017550	6 NM_008844	6 NM 008237	6 NM 033268	6 NM_015819 6 AK007800	6 NM_011156	6 NM 026593	6 NM_008743	6 M19413	6 NM 013636	6 AK014261	6 U69136	BC005502		7 NM 011750	7 AK015158	7 NM 010077		7 6003808	0.0

cluster analysis I breast cancor hypothalanus

Description SH3-domain binding prolein 5 (BTK-associated) tumor necross's factor abilish einduced prolein 3 neural precursor cell expressed developmentality down-regulated nedd8 kallukrein 4 (prostase, enamel mairix, prostate) g proferio coupted receptor 'amily' c group member h gpxc1h RIKEN CDNA 23 tool 41 17 gene creatine kinase, milochondrial 1, ubiquitous 18 days emviyor vitee rockat clone; 11 10004a08 RIKEN CDNA 28 105 10D14 gene phosphalidylinosilol glycan, class H	ornithine decarboxylase antizyme 2 RIKEN cDNA 1700018F24 gene RIKEN cDNA 1700008A04 gene RIKEN cDNA 2700097009 gene Similar to HCDI protein 184,44% Homo sapkens] zinc finger protein 1 Semilar to CDC 16 protein - human 78% N-methylpurine-DNA glycosylase offatory receptor mor 17-1	10 days neonale skin riken cdna done:4733401i23 glycoprotein-associated amino acid transporter y-tatta linked by disulfide bridge to 4f2hc/cd98; y+lattb glycoprotein-associated amino acid transporter y-tatta linked by disulfide bridge to 4f2hc/cd98; y+lattb dA2286 ER0-2-like protein_ELD-1 (98% human) a disinflegrin and metaloprotein-ass domain 19 (metim beta) a disinflegrin and metaloprotein-ass domain 19 (metim beta) grown lactor receiptor bound protein 7 chore mgc:5633; epidermal receptor-binding gb07 protein phosphalase 19 formetly 2c magnesium-dependent gamma isoform cone mgc:6666	serine/threonine kinase FKSG8144% human) Nyks pi Z probleh RikEN cDNA 170005608 gene phrosphoirlesterase related contactin associated protein 4 kandaptin RikEN cDNA 633057571 gene hypothetical protein FLJ13782 (94% human) anaphase-promoting complex subunit 7 copine III cathepsin d exon mesodern posicrion 2 mesp2	N.Z.Wdinerlyaguanosine fiNA metihyliransferase-like [Homo sapiens] 89 % RIKEN CDNA 4221508M14 gene SYO_HUMAN Gulaminy-IRNA synthetiase (GlutamineIRNA ligase) (GLNRS) 90.06 % SYO_HUMAN Gulaminy-IRNA synthetiase (GlutamineIRNA ligase) (GLNRS) 90.06 % In-acetylglucosanninyliransferase iii mgai3 glycosyliransferase introlless coding region: mannoside acetyl glucosanninyliransferase 3 I-IRNA MOLISE Histidine Iruda rucleotide-binding protein (Protein kinase C inhibitor 1) (Protein kinase 60 % small inducible cytokine sublamity cys.x-cys member 9 done mgc.6179. b scy69 Similar to KUP protein - human 89% RIKEN CDNA 49334 11 K20 gene A38346 Gulfe-high-suffur keralin 1 - mouse 32 % meiotic cohesion Rec8 c-type lectin dc11 type ii transmembrane protein RIKEN CDNA 1600012K10 gene
Locus Gene 24055 Sh3bp5 21929 Trilapi 18002 Nedde 56540 Kikd 1823 Gpcr1h 76946 2074 Spr1b 66492 110417 Pigh	18247 Oazz 69396 69351 7.2658 219094 22640 Zip1 7.2610 17477 Mpg 209102 mor17.1	Unknown 20540 Sic7a7 22561 Zipm1 66913 11422 Adam19 1698 Libp3 14786 Grb7 14208 Ppm1g	71099 70882 Nyd-sp12 70556 19212 Prer 170571 Cninapd 20534 Sid-alap 252973 GNIZ 56317 Apc7 70568 Cpne3 13033 Clsd	98685 99541 17094 Mgni3 68917 17329 Scy09 72788 66756 71648 8675 GCII 67009
Churdus Access 7 NM, 01894 7 NM, 009397 7 NM, 009897 7 NM, 005894 7 NM, 005894 7 AVANGES	/ MAKOSBS / MAKOSBS / MAKOSS / MAKOS /	7 AK014638 7 NM 014658 7 NM 019569 7 NM 025641 7 NM 005570 7 NM 010346 7 NM 010346 7 NM 010346	S 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	7 AK004833 7 AK004814 7 AK004814 7 AK004814 7 NM 010795 7 NM 008599 7 BC004761 7 AK013374 7 NM 025747 7 NM 025002 7 NM 019791 7 AK005037 7 NM 019791 7 AK005037 7 NM 019791

cluster analysis I breast cancer hypothalanus

Description galanin receptor 2 garana-aminobulyric acid (GABA-A) receptor, subunit gamme 1 hook nonclog 2 (Drosophila) G-protlein coupled receptor 12 argmase type II A56716 arromalic ester hydrolase (EC 3.1.1) - human 85 % lecithin-telinol acylitansferase (phosphalidy/choline-retinol-O-acyliransferase)	cell division cycle 42 homotog (S. cerevisiae) myosin heavy chain 3 flank mnyosin heavy chain 3 flank RIKEN cDNA 1700051CD3 gene RIKEN cDNA 1700057A23 gene RIKEN cDNA 5830412809 [Mus musculus] nsp-ike 1 protein nsp1 tms-sec and fosb: aa 1-338 nsp-ike 1 protein nsp1 tms-sec and fosb: aa 1-338	olfactory receptor 49 olfr49 limb deformity formin tifful frefoil factor 3/intestinal protein exons 1-3 GTPase regulator associated with the focal adhesion kinase pp (25 [Homo sapiens] 56.19 % GTPase regulator associated with the focal adhesion kinase pp (125 [Homo sapiens] 56.19 % Similar to MOUSE Alpha-actinin 3 (Alpha actinin skeletal muscle isoform 3) (F-actin cross linking protein) 28%	solute carrier family 30 zinc transporter member sk.30a3; znt-3 glycoprotein iib gpatpha ib-iig2b allete platetef fibrinogen receptor NYC2 transcription factor related, bocus 5 (Orosophila) aquaporin 7 sorting nexin 4 (Mus muscudus) 100 %	antigen p97 (metanoma associated) identified by monoclonal antibodies 133.2 and 96.5 small GPT (metanoma associated) identified by monoclonal antibodies 133.2 and 96.5 small GPT as home of Serevisiae) RIKEN cDNA 2700016508 gene RIYH3_RAT Myosin heavy chain, last skeletal muscle, embryonic (22% M.musoulus) vincutin syntaxin binding protein 2 RIKEN cDNA 3230401103 gene	persephin ras oncogene famity recombining binding protein suppressor of hairless-like (Drosophila) recombining binding protein suppressor of hairless-like (Drosophila) hypothetical protein FLJ20668 [Homo sopiens] 83.50 % zinc finger protein 119 - gonadotropin inducible
Locus Gene 14428 Gait 2 14405 Gabrg 1 14033 Hook 2 11847 Ag2 68021 79235 Lrat	12723 Clcn 1 12540 Cdc42 17833 Myn3 20024 Rpo2lc 1 88107 73975 76420 14912 Nhx6-2 68310 14282 Fosb	18348 Offr49 14260 Fmn 2.1786 TH3 71544 75356	22784 Sic30a3 16399 liga2b 16391 Nira2-5 11832 Aqp7 69150 Sira4	30060 Mit2 54170 Git2 67120 73332 22330 Vct 20911 Stubp2	71900 19197 Pspn 19138 Reb13b 19668 Rbpsuhl 71779 104349 Z[p119
Chuster Access 7 NM 010254 7 AF 156490 7 BC002226 7 NM 008151 7 NM 009765 7 NM 007557 7 NM 007557 7 NM 007557	7 AJ011106 7 AA018573 7 NM, 009881 7 M74753 7 MM,011294 7 AK015245 7 AK015324 7 AK015517 7 L08074 8 MM, 026670 8 MM, 026670 8 MM, 026670	8 NM_008163 8 AK012922 8 NM_019922 8 NM_019960 8 BC0065292 8 BC006526 8 NM_011575 8 AK018520 8 NM_013891 8 NM_013891 8 NM_013891		NM_013900 NM_017475 AK016428 AK006568 NM_016857 NM_016857 NM_01503 AK014327	8 NACOUSCAGE 8 NM_016858 8 NM_016858 8 NM_031494 8 NM_03186 8 NM_009789 8 NM_009789 8 NM_009789 8 NM_009789 8 NM_009789 8 NM_023020

FIGURE 20.13

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ysosomul pipha-ghucoadase Sumita to fyor I transmembrane treaper (setture-related protein) [37% Hono sapiens) Simita to fyor I transmembrane cepacit (setture-related protein) [37% Hono sapiens) Simita to fyor I transmembrane chare. (10001113 Selfytmischesse / (lighta-Al-ad-phreutaminy 2.3-betgablockeyf 1,3)-N-acetyl galactosammide alpha-2.6-dolytransferase) F Interest of the cecanic of the common recepacity of the common recepacity of the cecanic of the common recepacity of the cecanic of the common receptor and the cecanic of the cecanic o	24063 Spry1 50935 Spry1 50935 Spry1 50935 Spry1 50935 Spry1 50935 Spry1 50935 Spry1 50936 Try10 54538 Msrn 56538 Msrn 56538 Msrn 56538 Msrn 56538 Msrn 56538 Spryn 17688 Msrn 56538 Spryn 1769 Spryn 17759 1
garmia suduni or ine galda-alphia receptor garmia suduni or ine galda-alphia receptor nuccia - anno anno anno anno anno anno anno a	22319 Vamp3
RNA binding mall protein 6 carboxyl ester lipase gamma subunit of the gaba-alpha feceptor	19654 Rbm6 12613 Cel 14407 Gabrg3
RIKE'N cDNA 2010004P11 gene vorreconssal 2, receptor, 4 RNA binding mall potein 6 carboxy esset ligase camboxy esset digase	67889 22310 V2r4 19654 Rbm6 12613 Cel 14407 Gabra3
vomeronasal 2. receptor, 4 RNA binding multi protein 6	22310 V2r4 19654 Rbm6
Bookin Trate, Shrift an tresture free Control Cholosa 1 a califogen-responsive finger protein; done mgc:5886 RIKE'n cDNA 2010004P11 gene	22660 Trim25 67889
protein regulator of cytotrinesis 1-tike artult male small intestine riteen chtra chtra chtra (2010/074 1)	233406
inlegrin bela-6 subunit integral membrane protein	16420 ligb6
candidate laste receptor 11/2 g protein coupled internal monthrone protein	83770 Tas1r2
small inducible cytokine 828 scya28	56838 Scya28
100335 hypothetical protein KIAA0564 • human (fragment)(89% human)	71758
similar to S. cerevisiae RER1 [Homo saptens] 96 %	67830
intranuclear protein	16328 Inmp
inorganic evrephosobalase (Homo saplens) 71 %	74776
clathrin-associated protein 19 ap 19	11769 Ap1s1
small unique nuclear receptor co-repressor sun-cor corepressor for hormone receptors c1d; dna-binding protein c1d	57316 C1d
GNAS (quantité nucleotide binding proteto abbta stimutation) complex focus	14683 Gnas
coronin-3	23/90 Corolc
syndecan 1	20969 Sdc1
brain cana clone minch-3986 unnamed protein product ribosomal protein L.3-like	66211 Rpl31
	20724 SerpinbS
	13618 Ednrb
moesin Denotishmal bitoenesis (actor 3	17698 Msn 56535 Pex3
nuclear pore membrane glycoprolein 210 pom210	54563 Nup210
RIKEN cDNA 4930557821 gene	75308
toli-tike receptor 4 tumor suppressor region 10	21898 104 56368 Tsp10
	71000
Skityfransferese 7 (teipha-N-acetymeuraminy/ 2,3-betagaloctosyl-1,3)-N-acetyf galactosaminide alpha-2,6-sibyfransferese, hvoothetical 43,2 Kd moteio 145%. Hanny caniensi	50935 State
	50935 Stat71
	66329 21854 Timm 17
sprouty homolog 1 (Drosophila)	24063 Spry1
lysosomal alpha-glucosidase	14387 Gaa

FIGURE 20-14

Oveription RIKEN CDNA 4922442K08 [Mus musculus] 100 % RIKEN CDNA 231001 IE08 gene Opioid receptor, delta 1 rab27a member ras oncogene family ctone mgc:11677; es cells riken cdna ctone:2410003m20 aduli male lestis riken cdna ctone:1700001002	aduli male lesiis riten odna done:4930544110 PD2 and LIM domain 3 Iaminin-2 alpha2 chair m-chair merosin chair m-chair steroidogenic acule regulatory protein related ctone mgc:6996	pancrealic polypeptide RIKEN cDNA 2010003D20 gene suppressor of variegation 3-9 homotog 1 (Drosophila) RIKEN cDNA 2000018L1 gene RIKEN cDNA 200014C22 gene deoxyribonuclease 1 viirin likely ortholog of yeast ARV1 [Homo sapiens] 80 %	organic calionic transporter-like oxcli i expressed sequence R74862 kremen kingle-containing transmembrane protein kremen kingle-containing transmembrane protein orphan g protein-coupled receptor gpx 33 related to chemoaltractant receptors fibroblast growth factor 8 Mus musculus adult male testis cDNA, RIKEN full-length enriched fibrary, chone:4930525F21 product:hypothetical protein testicular cell adhesion molecule 1 protein phosphatase 1, regulatory (inhibitor) subunit 11	mucolipin 2 - RIKEN cDNA 3300002C04 [Mus muscafus] 100 % melanocortin receptor calpain 12 RIKEN cDNA 231004309 gene paternally eurossed 10 Smilar to CAMP [7 1% Mus muscafus] 2 103, HUMAND Zen finger protein 183 92 % transmembrane protein 4 S-hydroxytyptamine serotomin receptor 1a htr18 laterin lyn	10 day old male pancreas riken cdna done:1810061i13; c-lype lectin-like receptor 2 dec2 adult male testis riken cdna clone:1700122011 gbutamate receptor tonotropic ampa4 alpha 4 gria4 asporin precursor aspn preproprotein type i extracellular matrix leuchne-rich repeat protein hos:14 protein T17239 hypothetical protein DKFZp4348027.1 - human (tragment) 85 %	Similar to Ran binding protein 11 (IPO11) [95% Homo sapiens] RIKEN cDNA 4933440J22 gene iap homolog a mina apoptosis inhibitor of bacutovirus protein; bacutoviral repeal-containing 4 birc4 backed itament structural protein in tens.CP94 pellino 2 single-rrinded 2
Locus Gene 67544 69602 18386 Oprd 1 11891 Rab27a 75435	75212 53318 Pdim3 16773 Lama2 59045 Min64	19064 Ppy 72058 20937 Suv39H1 71831 72135 13419 Dnase1 74199 Vil 68865	18399 Sic22a6 97423 84035 Kremen 14762 Gpr33 14179 Fgf8 75823 75870 Tcam1	68279 1720 Mc3r 60954 Capn12 69671 66536 6638 6638 15550 Tmm4 15550 Hr 1a	56760 Clec.2 78651 14802 Giad 66895 Aspn 15401 Hoxad 76459	76582 74470 11798 Birce 1205 Birse 93834 Peli2 20465 Sim2
Churter Access  B NM, 024203  B AK009293  B NM, 013622  B NM, 023635  B AK005616  B AF 140708  B AK007936	8 AK016037 8 NM_016798 8 U12147 8 NM_025960 8 NM_025960	8 NM, 010164 8 NM, 00891 9 AK008060 9 NM, 011514 9 AK005893 9 AK01708 9 AK013183 9 AK004388	9 BC004098 9 MM 008766 9 RM 008766 9 MM 23296 9 X76011 9 AF045766 9 U18746 9 AR015893 9 AR015873 9 AR015973	9 MM, 02656 9 MM, 02656 9 MM, 00656 9 MM, 00656 9 AK00979 9 AK09979 9 AK014919 9 AK014919 9 NM, 008308 9 NM, 008308 9 NM, 008308	9 NM_01998 9 NM_01994 9 AK007241 9 AK007241 9 NM_019691 9 AK10695 9 AK13538 9 AK13538 9 AK019316	AKO 103.10 AKO 17.134 NM_00975.1 NM_033602 NM_011377 BCC0333.1

Description inferteron alpha formly, gene 4 crystalin appeal a milochondrial crystalin appeal and the formula control of the certyme 2. NAD(+)-dependent, milochondrial formula associated protein 3.1 [87% Homo sapiens] Y174_HUMAN Hypothetical protein KIAA0174 95 % hes2 junctophilin 1	merubrane-spanning 4-domains, subtamily A. member 6D. MS4A6D protein [Mus musculus] 100 %  CAMP [Mus musculus] 100 %  reserrchyme forth head-1 protein; mfh-1  SEC22 vaside traffiching protein; life 1 (S. cerevisiae)  telomerase reverse francopialise mfert, calabytic subumil polassium vollage gated channet subfamily in lagareled member 3 kcm3	nuclear receptor subtamily 5 group a member 2 nr5a2 Similar to ENP1_HUMAN Ectorucleoside triphosphale diphosphothydrolase 1 (NTPDase1) (Ecto-ATP diphosphothydrolase) 44% adult male lestis riken cdna clone:4822502b01 d-glycosaminy/ 3-o-sulforansferase-3b 3-ost-3b potassium channel, subfamily V, member 1	growth factor receptor bound protein grb 10 ribonucleotide reductase M2 KIAAD773 gene product [31,09% Homo sapiens]	And 0593 serminal vesicle secretary protein IV precursor - mouse (fragment) 100 % antigen identified by monoclonal entitody MRC OX-2 recepto reprosonate by monoclonal entitody MRC OX-2 recepto reprosonate by monoclonal entitody MRC OX-2 recepto reprosonate by monoclonal subtaining 4 domains, subfamily A, member 6C preprosonate by monoclonal 2 Similar to nuclear VCP-tike; Nuclear valosin-containing protein-tike (Homo sapiens) 87 % capial 12 in this one and protein tike from the protein tike from the protein tike 1 in this one territy member done mgc:6430; golgi snare 9527 secz2 vesicle traffiching protein-tike s, cerevisiae clane mgc:6437; golgi snare 9527 accyt-Coenzyme A oxidase 3, pristanoyt; pristanoyt	pleckstrin formology. Sec7 and collectool domains 4 wingless tellated MMTV integration site 8A dynamin calines in Zinc linger protein 275 putainver potein (28% Arabidopsis thaliana) mitochondrial solute carrier protein relinablastoma bindring protein done mgc.6013 riten cdna 4432404a22rik
15967 (Ina4 15967 (Ina4 12955 Civab 107029 Me2 69473 71955 15206 Hes2 87339 Jph I	68774 Ms4860 170717 14234 Foxc2 20333 Sec211 21752 Tert 16512 Komb3	26424 Nr5a2 72090 70954 54710 Hs3sl3b 67498 Kcnv1	14783 Grb10 20135 Rim2 78408	70355 77781 Mox2r 73656 Ms486c 73656 Ms486c 77257 Mecp2 67599 Capn12 14958 H10 68559 Metap11 56494 Gos12	72318 Pscdd 20890 Wnl8a 12330 Canx 12330 Canx 27081 2[p275 74090 19648 Rbbp 19648 Rbbp
Guster Access 9 NM, 010504 9 NM, 010504 9 RC(004709 9 AK0091 19 9 AK0017364 9 NM, 000236 9 NM, 000236 9 NM, 000236	9 NW_026935 9 AK020193 9 AK016845 9 AK118384 9 NW_013519 9 AV01359 9 NW_009354 9 NW_009354	9 AND 16649 9 NA, 030676 10 AK008590 10 AK018521 10 NM, 030808 10 NM, 02449 10 NM, 033327 10 NM, 033327	10 ARGATAZO 10 AMC10345 10 MM_010345 10 MM_009104 10 MM_0197 18 10 MM_008857 10 MM_008867 10 MM_008867	10 MAST72 10 NM 009431 10 NM 001325 10 NM 002855 10 NM 008051 10 NM 008051 10 NM 008197 10 NM 008197 10 NM 008197 10 NM 008289 10 NM 008289 10 NM 008289	10 AKO10508 10 INI_009290 10 INI_009290 10 INI_007597 10 AKO1074 10 AKO102491 10 AKO10200 10 INI_005033

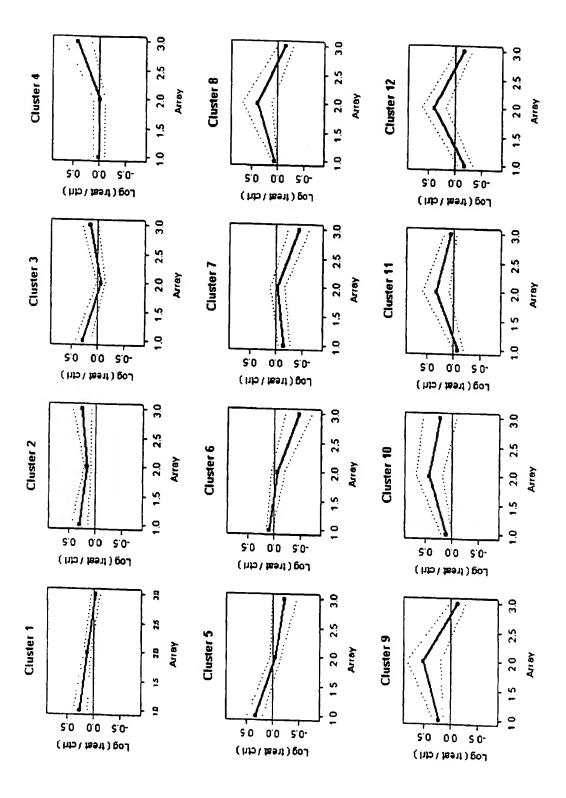
	2 IBIO TgDs (ransforming growth factor, bela induced 12547 Cded cell differentiation and embryonic development	57423 Atp5j2 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit 1, isoform 2 114896 Afg3i1 AFG3(ATPase family gene 3)-like 1 (yeast	67 i 12 Fg/22 fibrablest growth factor 22 16331 tpp actin-binding protein mipp	19 Lgi1 leucine-rich repeat LGI family, member 1 hypothetical protein FLJ 13253/7 1% furman)	Egr2	11782 Apds adult male cerebellum riten cona clone: 1500019b17		8 disintegrin and metaltoproteinase domain 12 metrin athra adam12	•	י עלונה . פונולותה חבום פג לוברתומת אישב לו תכנוג קלונה !	Galgi2 udp-n-acelyl-alpha-d-galactosamine:n-acelylneuraminyl-galactosyl-n-acelylglucosaminylpotypepiide-beta-14-n-acetylgalactosaminyltranslerase galgi2		1 Griebo 1 (grizocortoco modulator) element binding protein 3 (Mus muscutus) 100 % Grizoco 100 modulatori element binding protein 3 (Mus muscutus) 100 %	Akp3	UNIVORM - Interior of reaching to the control of th			Grypc garmina c-crystalin; garmina-c-crystalin	F	246	•		Can be Krycatherin Can be T 1946 and Can be T 19	Cst9	we i i a mework Economismo I i namong A i i. cereviside) E.ii enhance of zeste homoloo 2 (Drosconiia)	Alm alamin	6	Using 2 upregulated during skeletal muscle growth 2 Minos and an advantad deduced 2			Conm2 cyclin M2 divalent cation tolerant protein CUTA [Homo sapiens] 91 %
3				56839 Lgit 66039		-	•				14422 Gatgt2	0	11878 Arr	11648 Akp3	67490 108062 Cett2	14825 Gro		12966 Cryac	15374 Hnl	67467			12556 Cdn 16 68947	13013 Cst9	14056 Ezh2	280662 Alm	50518 @	83677 Usm	52575	53877 Ear4	94219 Cnnm2 67675
	10 NM_009369 10 NM_021280 10 NM_026320 10 NM_023547	10 AK005036 10 NW_020582 10 AK012394 10 AF375046	10 AK08922 10 AF285178 10 NM_009440 10 M63850	10 NM_020278 10 AK002441		10 NM_021710	10 AK009391 10 NM 025701	10 NM 007400						10 NM 007432	_	90	10 AK019788 10 AK017076	ş		10 NM 026177 6	NM_007411	AK003611	TO NIM UO/BB3 7,			••			•		10 NM_033569 94 10 AK002828 67

Obstription TDD automotion products to	John wanneshinding	ector, montoured, terky	CDKR_HUMAN Cell division protein kinase 8 (Protein kinase K35) 90 %		male germ celt-associated kinase	and the factor and and the factor an	Cell dimologic cycle 23 rightologic S. Lefensiae	arternin			arp) actin-related protein yeast homolog a centractin alpha clone mgc:5916; es cells riken cona clone:2410038/03	aop-rioosyii ansierasa nad - poy aop-rioose bolymetase adprtz; z parpz parp-1-tike protein parp-2				inhibitor of crowth family member 1-the soft above the market and 200 s.	neurolysin; neurolensin endoceolidase 190.33% Homo sapiens	mesendrymal stem cell protein DSC54 [92,03% Homo saptems]	Mus musculus adult male testis cDNA, RIKEN full-tength enriched (fibrary, clone:4930448A20 product:unclassifiable			DJ84_MOUSE OnaJ homolog subfamily 8 member 4 100 %	elk4 member of ets oncogene family done image:3589378	Calsyntenin-1 protein	melanocyte prolifeating gene 1		eci2 oncogene	histocompatibility 2. Oregon apple focus	addin mare result inten cone cone: 35,14,2001.3 Tul misen sequence; collagen B). XIX chain	es cells riken cana cione:2410018020	oliatory receptor mor 17-1	дурісэл 6 дрс6		gcn5 histone acetylicansferase; clone image;3491089			ACM3_MOUSE Integrating acety/choline receptor M3 (Mm3 mAChR) 100 % /	KINEN CUNA Z31000/H09 gene	Advisory Divient	advanced gyrosylation end product-specific receptor		mannosidase 2 alpha man2a1			halap-x haptoid specific alanine-rich acidic protein		Then come 1500011611 15000116118	DIVERSITY OF THE PROPERTY OF T	TANEN CORA 24 10 13 M 17 Bene			Cg10671 like (Drosophila)		Soulte Cartier (armly b neurotransmuller transporter Burnne member sicoad; mus cookii Baumedella-dianine Isrambrovia seriasiiva nana 1			
Locus Gene 62088 To 120h	70361 Lman 1	16469 Jrk	67587		17152 Mak	47677 (445)	יבאב במנגאנ	11876 Artn			54130 Actr1a	Zudow owe I I				69260 (no.1)	75805	71373	73993			67035	13714 ER4	65945 CsIn1	60315 Myg1	1	13605 Ect2	15001 H2-Oa	1861BO C7071	72083	209102 mor 17-1	23888 Gpc6		14534 Gcn5l2	77.77	11971 Algn1	12671 Chrm3	67677	11505 Anns	affw occi		17158 Man2a1			60600 Наварх	000000000000000000000000000000000000000	CEDED DATA	76820	^ 700 /			68680 Cg10671	0-0-0	21366 Sictate			
Chuster Access	10 AKD11495	10 NM 008415	10 AK015854	11 NM 025380	11 NM 008547	11 ABU49821		11 NM 009711	11 NM 025383	11 AK01/655	11 NM 016850	11 NM 026581	11 AKO12848	11 NM 0251	f 1 AKO 10807	11 NM 023503	11 BC016224	11 AK017282	11 AK015411	11 NM_025826	11 AK015825	11 AK008537	11 NM 007923	11 NM 023051	11 AK017955	11 AF145716	006/00 MN 11	11 NM 008206	11 NM 010590	11 AK010544	11 AF 133300	11 NM_011821	11 AKD05311	11 NM 020004	11 AK011747	005 /00 MM 11	11 NM 033269	11 ARGOS (0)	11 ACOTOON	11 NM 028119	11 BC005574	11 NM 008549	11 AK004004	11 AK008516	11 NM 021898	11 60005625	0200 MM 11	11 AKO10821	11 NM 013529	11 NM 019677	11 AK006136	11 AK003939	11 000 MN 11	11 NM 009320	11 NM 018768	11 NM_030253	11 NM_020267

Overripsken small nuclear RNA activating complex, potypeptide 3, 50kD [84.95% Homo sapiens]	CUSS. MOUSE Pulsive protein C2 tor158 homolog 54.41 %	frc. fringe-like 1 (Orosophila) RIKEN cDNA 2300002018 gene	1) nudix (nucleoside diphosphale finked molety X)-type molif 11	Friend virus susceptibility 1		44 nair kerain adok 5 na5 4bor transcription factor tbx 1	heterogeneous nuclear ribonucleoprolein c hampo	ate of anti-transmission and anti-transmissi		•	5-3 excribonuclesse 1	Gambiotike Sambiotical methyleneletrahydrototate dehydronenas-krathydrotase mitortyddiol mae'n 20 k. MIDC FlluMA Bifunctional methyleneletrahydrototate dehydronenas-krathydrotase mitortyddiol mae'n 20 k.	chymolypsin activity, growth factor activity	_	2	RIKEN cDNA 1700095KDB gene		Controlled revenue &		tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein sigma polypeptide ywhas; 14-3-3 mme t		plasmacytoma expressed transcript 2			cachem EGF LAG seven-pass G-type receptor 1; cacherin EGF LAG seven-pass G-type receptor (25% Mus musculus)	Similar to cytoplasmic dynein heavy chain 2 [98% Rattus norvegicus]		Similar to CGHU2V collagen alpha 2(V) chain precursor - human 30%	RIKEN cDNA 2610027F03 gene			eik i member of ets oncogene family	and a land a state of the second as a confidence of the second se	proxingiamum X, proxincycum symmetre prigs A38647 ghulammale-rich protein form A. submandibular gland - rai 31 % /
Locus Gene 77634	74281	76157 Freil 69541	58242 Nudit 1	14349 Fv1	74772	21380 Tbx1	15381 Hnrpc	14661 Ghid	18366 OlfreS	20502 Slc16a2	24127 Xrn1	68621		12151 Bmit 67285 Sdeepol0	170658 Ndufs5	74286		14785 Gorso		55948 Mkrn3		18530 Pet2			78249	77047		72125	69904			137 12 Elk 1	mirror coccus	71400
Chuster Access 11 AK019863 11 NM 023884 11 AK008838 11 AK017509	11 AK006994 11 AK015078	11 AK018094 11 AK009014 11 AK014338	11 AB041576 11 AK020583	11 NM 010244	11 AK003623 11 V00711	11 AF326960	11 NM 016884	11 AK019434 FF1900 WM 11	11 NM_013617	11 NM 009197	11 NM 011916	11 AK003824	11 X76290	11 M64067	11 BC002163	11 AK007086	11 AK016612	11 NM 010340	11 NM 020563	11 NM_018754	11 NM 024469	11 NM 025462	11 AK015743	11 NM 026428	11 AK019508	11 AK014500	11 AK011555		11 AK011565	11 NM_011973	11 NM_013732	11 NM_007922 11 NM_025736 11 AF220294	11 AK019591	

Description e21 tike transcriptional repressor protein ema 5.3-ructeotidase, milochondrial Similar to CO8G_HUMAN Complement component C8 gamma chain precursor 73 % human	neuregulin 3 hsp60 protein ctones 313-7 -9 and -m1 555 aa; cheat shock 65 thymine DNA glycosylasa solute carrier family 15 (H*/peptide transporter), member 2	Moloney leukemia virus 10-like 1 [Mus musculus] 37.37 KIAA0547 gene produci [80% Homo sapiens] collagenous repeal-containing sequence of 26kda protein cors26 putative secretory protocadherin 12 pcdh12	stometin-like it: stomatin-tike protein 1(79% human) prodactin-tike protein E fibrobasi gowth lactor receptor 4 fight a novel member of factor family von Ebner minor salivary gand protein transforming growth factor beta receptor 1	138191 nucleic acid binding protein • human (fragmen!) 98 % human keralin associaled protein B-1	parathyroid hormone precusor pth traf family member associated nt-kappa b activator tank; traf-interacting protein i-traf	erythrocyte protein band 4.1-like 4a inducible T-cell co-stimulator fidgetin-tike fight1 aaa family protein A31589 carboxypeptidase C (EC 3.4.16.5) precursor • human 39 %	Fas death domain-associaled prolein RIKEN CDNA 2610005M20 gene MAP/microlubule affinity-regulating kinase 2 milochondrial ribosomal protein t.15	neurotensin receptor prosome mecropain subunit alpha type 8 CG1-143 protein [Homo sapiens] 81 % / Mus musculus na indochondrich, complete genome Mus musculus 13 days errubyo mate testis CDNA, RIKEN full-length emiched library, done:50:30449J20 product:hypothetical protein ibosomal protein L37	Imp. 2 fissus inhibitor of metalloprotenases type; metalloprotenase GPI-anchroted HDL-thruting protein 1.  adult male cerebellum riken cdna clone; 15000320.24  rox protein protein chi basic heiki-loop-helix leucine zipper protein; myc antagonisi mnt max-network bhihzip adult male testis riken cdna clone; 1700025113; pyruvate dehydrogenase pdha-2 calcineurin B homologous protein 2  tulletin-interacting protein 39 tip.39	days neonale head riken cdna done:5430425H06; 9130005n14 9130005n14frik
Locus Gene 50496 E216 103850 NISm 69379	18183 Nrg3 15510 Hsp60 21665 Tdg 57738 Slc1592	71159 75894 81799 Cors 53601 Pedh12	69106 19113 Pripe 14186 Fgfr4 228801 21812 1gfbr1	66618 16703 Krlap8-1	19226 Pth 21353 Tank	13824 Epb4.149 54167 Icos 60530 Fign11 71287	13163 <i>Daxx</i> 70308 13728 <i>Mar</i> k2 27395 <i>Mrp</i> t15	18216 Nisr 26443 Psma6 69168 17722 ml·Nd6 68148 67281 Rpi37	2 1858 Timp2 68453 Gpibb1 18012 Neurod1 17428 Mni 18589 Petha2 70261 Chp2 54723 Tip39	68303
Chuster Accuss 11 NM_033270 11 AK008226 11 AK005962 11 AK012765 11 AK011765	11 NM_008734 11 X55023 11 NM_011561 11 NM_021301 11 NM_026119	11 NM_023625 11 AK016833 11 AK016299 11 NM_030888 11 NM_017378	11 AK007508 11 NM_008930 11 NM_008011 11 U46068	11 NM_009445 11 AK011937 11 D86423 11 NM_031396 11 AK020207 11 NM_019914 11 NM_031182		12 NM 024260 12 BC007166 12 NM 017480 12 NM 021891 12 AK018593 12 BC002251 12 AK017087	12 AK007781 12 NM_007829 12 NM_010016 12 NM_013556 12 AK011326 12 NM_007928 12 NM_007928	12 NM_018560 12 NM_01856 12 AK007718 12 AK020165 12 AK020067 12 AK020067	12 NM 011594 12 AK003305 12 AK005073 12 NM 010813 12 AK 008811 12 AK008392 12 AK020306 12 NM 018783	12 AK017385 12 NM_026667 12 AF291660

Description melanoma aningen, (amiy L. 2 adult male testis riken cdna clone:4930430k04	uramodulin CDK2 (cyclin-dependent kinase 2)-associated protein 1 ribosomal protein s29	RIKEN cDNA 84304 16G 17 gene lectin, galactose binding, soluble 9	rines Conv. Conv. Octobers 1. gara 145081 hypothetical protein C316G12.2 (imported) - 73% human Ihyroid hormone receptor interactor 10 similar to -pir:731100 - 731100 probable podassium channel 2 - rat 89 %	Gardner-Rasheed Ielne sercoma viral (Fgr) oncogene homolog los-related antigen-2 rrusclebtind-like (Drosophila) JC4131 glioma pelhogenesis-related protein - human 40 %	S12207 hypothetical protein (82 element) - mouse (68% human) interteon-gamma (muith-gamma) myosein-tike myozk myosein-tike myozk myosein-tike myozk septembane molien 4	programmed cell death pdcd i	or an expressor, associated with recovery thoughts specific chaperone of midnoish RIKEN cDNA 4930544L04 gene
Locus Gene 27385 Magel2 74638	22242 Umod 13445 Cdkap1 20090 Rps29	71469 16859 Lgats9	68327 106628 Trip10 77536	14191 Fgr 14284 Fosl2 56758 Mbrt 69286	71546 15978 IIng 59006 Myoz2 56214 Scampa	18566 Pdcd1 17527 Mpv17	108903 Tbcd 59090 Midn 78807
Chutes Access 12 NM_01379 12 AK015254 12 AK01580 12 AK008793 12 NM_013563	12 AK004852 12 AK004852 12 NM_009093	12 AK018420 12 NM_011651 12 NM_010708	12 AK021319 12 AK002309 12 AK021149	12 NM_01020B 12 NM_008037 12 NM_020007 12 AK005860 12 NM_026047	12 AK018561 12 K00083 12 AK015259 12 NM_021503 12 NM_019575	12 NM 008798 12 NM 027702 12 NM 026632 12 NM 008622	12 AK020685 12 X81632 12 NM 021565 12 L29479 12 AK019737



Description		hypothetical protein MGC 15435 [Hamo saptens] 51 %						_			_		Rabbinite action of the mention 23 35 %										DYNC_HUMAN Dynactin complex 50 kDa subunit (50 kDa dynein-associated polypeptide) (Dynamtin) (DCTN- 94 %	RIKEN CDNA 2610206C17 gene	_		Similar to APXL_HUMAN Apical-like protein (APXL protein) humna 28 %			hypothetical protein FL23467 (Homo septens) 93 %	microsomal inglycense items of protein	Ni protein 2 (89% Horno Sabens)	to days entaryoung tiken come contractions		DINEON CONTROL CONTROL DINEON CONTROL DINEON CONTROL C	Anne Common III and Anne C	Graphin I 100 November 100 November 1 100 November	trehalase (brush-border membrane grycoprotein)	reticutocalbin 2	hypathetical protein FLJ20080 (Hamo sapiens) 78.35 %	Similar to PRP1_HUMAN Sativary protine-rich protein precursor (Chones CP3, CP4 and CP5) (Basic peptid 35 % human	_	hypothetical protein MGC14827 (85% human)	RIKEN CDNA 2310016G11 gene	RIKEN CONA 4931431F19 pene	RIKEN CONA 1500035H01 pere	RIKEN CDNA 4930451F05 gene	A I Pase, class VI, type 11A	MUST STANDARD KINETON COUNTY AS 101 16 GOOD GROVE (2410) 16 GOOD KINETON COUNTY (AMERICAN COUNTY AS 101 16 GOOD GROVE (AMERICAN COUNTY AS 101 16 GOOD GROVE (AMERICAN COUNTY AS 101 16 GOOD GROVE (AMERICAN COUNTY AS 101 16 GOOD GROVE) AS 101 16 GOOD GROVE (AMERICAN COUNTY AS 101 16 GOOD GROVE (AMERICAN COUNTY AS 101 16 GOOD GROVE) AS 101 16 GOOD GROVE (AMERICAN COUNTY AS 101 16 GOOD GROVE (AMERICAN COUNTY AS 101 16 GOOD GROVE) AS 101 16 GOOD GROVE (AMERICAN COUNTY AS 101 16 GOOD GROVE	KINEN CITINA 31,3240/ CLS gene	GOUDE C. Detail	STREAM OF THE ATT SPORTS AND A CONTROL OF THE SPORTS OF THE TAKEN SPORTS OF THE SPORTS	arin hels sections			Similar to Ubiquitin-conjugating enzyme E2-23 kDa (Ubiquitin-protein ligase) [42% mouse].	sp P46096 - SYT1 _MOUSE Synapidiagmin I (Syt1) (p65) 38 %	culin 2	Fancori anemia, complementation group G	T 125 15 hypothetical protein DKF 2p434B103.1 · (28% human)
Gene			Parva	OII, 160	Hsna2	Brink	Sor	Msdal		NdufbS	5	; i		Rannta	420	}	CASS	Pdcd 10	NAbit	Gmfa	Pripm	Torib				Fap			Odz2		Q III			0000	6250	č	5	Treh	Rcn2			Mrps18a						A(D118		46-00	02300		ŧ	Offr 159				COIS	Fancg	
Locus	101513	72114	57342			-	20675	14126	75062	66046	81879	70502	74239	72297	20303	71712	23833	56426	18038	63986	56635	30934	69654	72495	87678	14089	71774	19607	23964	69/69	7771	52633	20000	6666	77616	57370	68876	58866	26611	216549	67998	68565	68201	69578	20980	76568	74673	07/06	2000	13447	73467	66440	11461	29849	56279	66799	77314	71745	60534	70892
Cluster Access	1 AK012535	1 AK002774	1 NM 020606	1 NM 030553	1 NM 008301	1 AF233580	1 NM 009237	1 NM 013516	1 8C009141	1 NM 025316	1 NM 023755	1 AKD 17530	1 AK006472	1 NM 028189	1 NM 013652	1 AK004552	1 NM 013706	1 NM 019745	1 NM 010909	1 NM 022024	1 NM 019991	1 AJ297743	1 AK009749	1 AK011897	1 NM_026309	1 NM 007986	2 AK004934	2 AF358257	2 NM_011856	2 AK007540	Z NM 008642	Z NM 023175			2 AKO21182	2 NM 020595		2 NM 021481	2 NM 011992	2 8C004630	2 AK004960	Z NM 026768	2 NM_026613	2 AK009387	2 AK016497	2 NM 023831	2 AK015385	2 NM_015804	Z NIM UZB630	2 MAC MOTOTO	2 AM 00/8/3		2 NW 007393	2 NM 019476	2 NM 019833	2 AK015348	3 AK021056	3 AK016520	3 NM_053081	3 AK014905

Gene Description YA02_HUMAN HYPOTHETICAL PROTEIN DJ1198H6.2 (50% HUMAN) Chordct Cysteins and histolinearch domain (CHORB)-containing, zinc-binding protein 1 DNA segment, Chr. 7. Roswell Park 2 complex, expressed Cpb binding protein FLJ1348 (Homo sapiens) But the phypothetical physical phypothetical p	Sca2 spinocerebellar alaxia 2 homotog (human) Similar to zinc finger, BED domain containing 3 (2BED3) [51% Homo sapiens] 21 (3939A elongin 8 [Homo sapiens] 72 % Fh1 furnated hydalases 1 capacity (antibilar) subunit 3 f [83% Homo sapiens] colactor required to So1 transcriptional ectivation subunit 3 f [83% Homo sapiens] protein phosphalase 1, regulatory (inhibitor) subunit 3 f [83% Homo sapiens] Grirpx guanine nucleotide releasing protein x Psma3 proteasome (prosome, macropain) subunit, alpha type 3 Trim 11 intpartite most protein tim 11 Trim 11 Intpartite most protein tim 11 CCD bit MAND Probable numitional contracts of the protein proteins of the proteins of th	Scriba sodium channet, voltage-gailed, type IX, aphia polypeptide Cash te casein kinase 1. epsilon Case (Cat-ba more). Voltage-gailed, type IX, aphia polypeptide Casein kinase 1. epsilon Case (CAt-ba more). Voltage-gailed, type IX, aphia polypeptide Casein kinase 1. epsilon Casein kinase 1. epsilon Casein kinase 1. epsilon JCGS4T high sulfur protein BZE - rai 37 % Mouse 6 days necnal ehad cDNA, RIKEN tull-ength enriched tibrary, clona:5430417C01 product hypothetical mucin 10, submandibudar gand salivery mucin [Mus musculus] 30 % RIKEN cDNA 170006311 gene RIKEN cDNA 4930556324 gene Clog companibility 2. M region locus 1. q subcomponent, ganna polypeptide Clog complement component 1, q subcomponent, ganna polypeptide 90306071.77Rix RIKEN cDNA 903056071.7 gene		
Locus 77632 66917 74322 67876 73634	20239 S 72114 72391 14194 H 14194 H 14199			
Access AK019795 NM 025844 NW 037080 AK010337 AK005551 AK007415		NM_018852 73 NM_018852 73 NM_018852 73 NM_018665 72 NM_018666 73 AK001866 73 AK001866 73 AK001868 73 AK001868 73 AK001864 74 NM_010899 16 NM_001854 17	THE PROPERTY OF THE PROPERTY O	5 NIM_009757 121525 5 AF067063 16001 5 AK0 10800 76791 5 AK0 10800 76791 5 NIM_009255 20720 5 NIM_009275 12425 5 NIM_00927 12425 5 NIM_009
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Gerre Description		-	Poid peptidyforalyl isomerase D (cyclophilin D)		Pipn12 pratein tyrosine phosphatase, non-receptor type 12	Nago N-acety galactosaminidase, alpha	Mouse 4 days neonate mate adipose cDNA, RIXEN full-length enrithed library, clone: 8430316H17 product through their	Usp14 ubiquitin specific prolease 14	_		Ornal darlin marks extens 1			Sylvaba sylvaban property			۰	Official oraciony receptor 154		Pgs 1 phosphalidydycerophosphale synthase 1		•	T-cell receptor beta, variable V20	hypothetical protein, 154			=	Blk B lymphoid kinase			Prss11 protease, serine, 11 (lgf binding)		m		Olife? Olife? oliectory receptor 67	Similar to mammalian inositol hexakisphosphate kinase 2 [Homo sapiens] 91 %				_	Chk 1 chaline kinase-like	Tetraspan NET-6 (95% human)	Cidn4 claudin 4	Abca 1 ATP-binding cassette, sub-family A (ABC1), member 1	Ī	Octa2eo5 CDC42 effector protein (Rho GTPase binding) 5		pir 142647 - 143647 hypothetical protein OKEZ 6434A2115.1 - himan transmit 92 %	printed and comments Addition	Ehd Ethemion records		Colds altha reliance in relati	fxyd? FXYD domain-containing fon transport regulator?	_	_	Gabrb 2 gamma-aminobulyric acid gaba-a receptor subunit beta 2 gabrb 2	Potr2g polymerase (RNA) II (DNA directed) polypeptide G	hypothetical protein FLJ23045 [26% Homo sapiens]	hypothetical protein DKFZpS86E1923 [Homo sapiens] 88 %	RIKEN CDNA 3110041018 gene	Onajb7 DnaJ (Hsp40) homolog, subfamily 8, member 7	
1960	12466	14836	67738	78052	19248	17939	78687	59025	23880	72181	13406	CABA		22473	2000	20000	436434	2/216	69885	74451	17688	12563	21594	57428	67963	93861	14211	12143	26447	57267	56213	72014	13525	18992	18368	16500	11989	67245	18546	67181	12651	68109	12740	11303	68735			68259	52504	57440	67608		57780		_	14401		74478	66291	73166	57755 C	
		Ž	6 NM 026352	6 AR01980/	6 NM 011203	6 NM_008669	6 AK021025	6 NM_021522	6 AF061744	6 AK009285	6 AJ242625	6 NM 017584	S MAI OTTEOR	e Aktocose	6 MM 010409	DESCRIPTION OF	6 AF 262302	97/510 WN 9	6 AKOTIZ/9	6 AK011463	6 NM_010830	999200 WW 9	6 X59150	6 NM 020555	6 NM_023409	6 NM 033149	6 NM 008017	6 NM_007549	6 NM_011972	6 NM 018758	6 NM 019564	6 AK005160	6 AF 167404	6 NM 008899	7 WM 013619	/ AK005166	7 AK010322	7 AK020915	7 NM 008791	7 AK012063	7 NM_007692	7 AK012571	7 NM_009903	7 NM 013454	7 AK004139	7 NM 021454	7 AK004164	7 AK019542	7 AK018444	7 NM 020578	7 AK013432	7 NM 007734	_	7 AK014490	7 NM_011069	7 NM_008070	<u>۔</u>		_	_	B AK006201	

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calpain 10
hypothetical protein MGC10771 [78% Homo saplens]
RIKEN ALM 64 100 11 D08 gene
Man muscuus acul nate testis cDNA, RIKEN full-length enriched förlary, clone:4933409K03 product.hypothetical Heat shock protein hsp70/Ankyrin-repeat/Yeast DNA-binding domain cyclic nucleoide gated channel cncg4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         leroid 5 alpha-reductase 2-like: HSAR gene; steroid 5 alpha-reductase 2 like (Mus musculus) 100 % placental protein 11 related signal recognition particle 14 neuronal pentrain 2 rights; np2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rec (Hono sapiens) 60 %
3 (1015) transforming protein llm - 56% mouse (strain balb/c)
1- box and teacher-rich repeat protein 12
1- box and electine-rich repeat protein 12
1- box super-lamby super-lamby protein lusy
1- box super-lamby protein lusy
1- box super-lamby protein lusy
1- box spatial protein lusy
1- box super-lamby protein lusy
1- box super-lamby protein lusy
1- box super-lamby super-lamby super-lamby protein a super-lamby su
                                                                                                                                                                                                                                                                                                                    widely-interspaced zinc finger motils
RikEN cDNA 170025602 gene
nasopharynges petineline specific protein 1
piccolo (resynaptic cyonnatrix protein)
vascular Rab-GAP/TBC-containing: BUB2-kire protein 1 [Mus musculus] 48.54 %
altohno dehydrogenase 6 (dass V), pseudogene 1
myosin binding protein C, cardiac
CD24e entigen
                                                                                                                              nebulin-related anchoring protein
BRF2, submirl of RNA polymerase III transcription initiation factor, BRF1-tike
BRF2, BDNA 4390339A66 gene
protease (prosome, macropain) 26S subunit, ATPose 1
protease (protein 4250 18-1)
   Description
adul male Iestis riken cdna done: 1700048h20; clone: 4933432e16
Iestis expressed gene 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cycloric T-hymphocyte-associated protein 4 junction-medialing and regulatory protein toll-interleukin 1 receptor (TIR) domain-containing adaptor protein NADH detivingenase (ubiquinone) 1, subcomplex unknown, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T09013 RING finger protein Fxy - 24% mouse
Similar to hypothetical protein FLJ10587 [Homo sepiens] 98 %
G protein-coupled receptor 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       panceas specific transcription factor, 1a
RIKEN CDNA 920528E23 gene
Phospholipase C, della
BRCAL associate RING domain 1
hypothetical protein FL23119 (65% human)
phosphatidy/inositol 4, phosphate S-kinase, hype 1 beta
                                                                                                professome (prosome, mecropain) subunit, beta type 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein phosphalase 2a. catalytic subunit, bela isoform
elektryctic itrasfation initiation factor 3, subunit 4 (della)
synaptic nuclear envelope 1
epithetial membrane protein-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ribosomal protein S6 kinase
RIKEN cDNA 5730416020 gene
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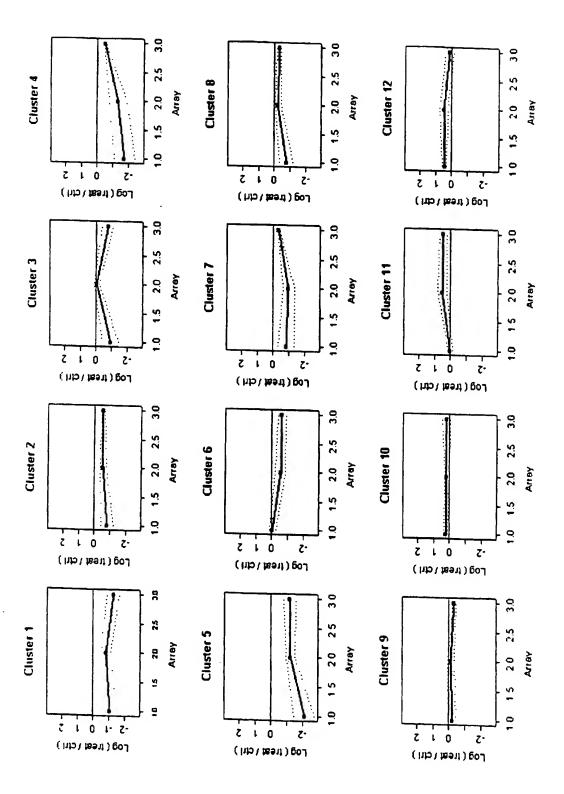
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Christer Access	10 NM 030744	10 NM 016693	10 AKO 16900	10 NM 007903	10 BC014706	10 AK017036	10 NM 009453	10 NM 029199	10 NM 010482	11 NM 019831	11 NM 007939	11 BC004794	11 NM 009660	11 NM_018755	11 NM 019800	11 NM 025978	11 NM 008782	11 AK015245	11 NM 021351	11 NM 025319	11 NM 009177	11 NM_008956	11 517857	9011111	12 NM 015260	12 NM 009881	00000 MM C1	12 AKO02722	12 NM 013618	12 NM 080428	12 AK005213	12 AK004795	12 AK015647	12 NM 030113	12 AK013274	12 NM 008716	12 AKON351	12 NM 052977	12 NM 011500	12 NM 009992	12 NM_019964	12 NM 024225	12 AK020538	12 NM_016926	12 AK019809	12 NM 00/462	12 AF 2B1045	12 NM 010883	25 MM 010053	12 NM 022030	12 AKO08753	



Gene Description Xist inxictive X specific transcripts, gene with no protein product			refrNP_055672.1 - KIAA0471 gene product (Homo sapiens) 93.55 % Sreb3 SREB3			Abio I ankyrin repeat and BTB (POZ) domain containing t Mvot metanocyte proliteating gene 1				•	Scn5a sodium channel, voltage-gated, type V, alpha polypeptide	FINEN CLOVE I COVER I LOUGH STATE I COVER I CO		235 ztp-35 zinc linger protein zfp-35 aa 1-580; exon		f Kninsilike farter 1 fandhmidt				(2 serum/glucoconticoid regulated kinase 2	and 1 putative N6-DNA-methydransferase	Moloney leukemia virus 10-like 1 (Mus musculus) 37.37	leucline-rich and death domain containing, p53 protein induced, with death domain [Mus musculus] 40.00 %	RIKEN cDNA 5830462121 gene	at hoved						is only touristicated protein yeass monstood a centracum alpha conte myc.30 for es cens man conte contes.24 tou.30 p.s. novdocin_manmonhemis							opioid growth tactor receptor				integrin alpha e epithetial-associated (tgae	small nuclear RNA activating complex, polypeptide 3, 50kD 184-95% Homo sapiens)	
	14065 F243 18806 Pld2 27374	22325 Vav2			37 FdR1	60315 Mva1	24071 Synj2bp					Norm	i S	22694 Zfp35		K K		2 Hexb		9 Sgk2	8 NSamt1	6	ю.	ın.	Haya3			Kcnq5		Nrahe	č	Polr2		Gosr2		Pes 1	,	<u></u>				ltgae		
Locus 213742			70771	21961	14137	60315	240			12540	20271	83485	75036	2269		16596	12111	15212	74264	27219	67768	71159	70388	46106	15400	83434		77887	13211	20186	18429	69920		56494	70804	64934		5/02/				16407 ltgae	77834	
Cluster Access 1 X59289 1 NM 026161	1 NM_007975 1 NM_008876 1 AF1G7573	1 NM 009167	1 AK014326	1 NM 027884	1 NM_010191	1 AK017955	1 NM 025292	1 NM_026047 1 AK016963	1 AK003359	1 NM_009861	1 NM_021544	1 NM 031375	1 AK015642	1 NM_011755	1 AK003808	1 NM 010635	1 NM 007542	1 NM 010422	1 NM_028842	1 NM_013731 1 AK014579	1 AK012019	1 AK016833	1 AK008713	1 AK018027 1 NM 013539	1 711717	1 NM 031255	1 BC004027	1 AK020325	1 091922	801600 MM 1	1 NM 011025	1 AK012635	1 AK005535	1 NM_019650	1 AK014543	1 NM 022889	1 AK019884	1 NM_0313/3	1 NM 010771	1 1.26164	1 AJ288061	1 NM_008399	1 AK019863	

Cluster Access	Locus	Gene	Description
1 AK014447			
4 AK007145			
1 AKO16162			
1 AK005564			mus musculus aduli male testis CUNA, KIKEN Millength enriched library, clone:4930557G23 product:homeodomain interacting protein kinase t
1 BC011457			
1 NM_019664	16516	16516 Kcnj15	potassium inwardly-recitying channel subfamily i member 15 kmil 15 imwardly re-ritkino k+ kird 2-a
1 AK020687			Barton Addition (Assessment Section )
1 NM_019445	54418	Fmn2	formin 2
1 NM_023066	65973 Asph	Asph	aspartate-beta-hydroxylase asph; aspanyl beta-hydroxylase
1 NM_009091	20054	Rps15	ribosomal protein s15 rps15
1 NM_019481		Slc13a1	solute carrier family 13 sodium/sulphate symponters member skc13a1
1 AK010475	69740		CGI-30 pratein [Homo sapiens] 85 % /
1 NM_011478			
1 NM_011825	23893 Prdc	Prdc	protein related to DAN and cerberus
1 NM_021456			
1 AND 10430	,		
1 AKOO5730	18534 PCK1	Š	phosphoenalpynuvate carboxykinase 1, cytosolic
1 NM 007385	41476 Appen	Anna	VII. 2 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 -
4 AKOOAEE	6761	Apoce	
4 AK013383	77550		Similar to Hir Par Groten 1978 Human
1 AKOO6318	8007		KINEN CUNA ZVODOZNOG BENB
1 AKOUSS 10			
1 MM 023554			
1 01/006169			
1 582853			
1 AIR4 007433	0.0040	0.00	
1 MM 015801	50757 AUG	AKDS	elikaline prospriatave intestine not mni requiring axps
4 44 02200	200	107	
266220 WW .	65106 Antigo	Antoipa	ADY-mosylation-rike factor 6 interacting protein 5
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1 MM 000 170	m6ds 50171	mgda.	esemble mulase
1 AKOO271	14047	1480	normeobox gsn-1
1 NM 007825	13123 Cvn2h1	Vn7h1	10.14 days sembors also ados alors of the contract of the cont
1 AK015001	66733		Control of the Contro
1 AK016509	2		Smind to potassium vollege-galeu channer, suorannet C, member 1, potassium channer RMZ (78% Homo saprens)
1 NM 023637			
1 AK013872	72797		DNA hinding modif amother (2) and adding henries an unbande, temporal and the contract of
1 NIN 007770	20.00	1	The change from forem 12, putative oran madeany-targeted protein 155.09% mono sapiens
* NM 025633	66550 Motost	ist if	c-ints proto-drougene protein precursor as -19 to 957; adult male liver riken consideration.
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1 N/M 045643	20043	d Silver	carnage originates many protein (mus muscalus) 100 %
1 NIM OCCURA	20932 SUN	Some Took	Surface 4
• 000000	27040	פנום	
1 AKO13281	7/6/0		looses I adenosma inprosphatose(99% numan)
1 AK016542			
1 NM 025911	67015		RIKEN EDNA 1810/60 ID IM-se muscanise)
2 MAA 024472	70554		TABLE OF CONTROL OF CO
2 NM 010004	T CO		pularive grycoipio uransier proteim (25% numan)
2 NM 008406			
2 NM 008072			
2 NM 008471	16669 Krt.19		aratin 10
2 BC004635	102607		SINX.1 HUMAN Sorting nexin 19 (84% human)
2 AK014112			
2 NM_023191			
2 NM 007527	12028 Bax	¥	10 11 days embryo riken cdna clone:2810443m09; bcl2-associated x protein bax
		ţ	

cluster analysis t lung cancer hypothalamus

Gene	2004			3983 Esr2 estrogen receptor beta estra entre entre					_	18131 Notch 3 Notch gene homolog 3 (Drosophia)	of ND 115187 1 - hynothetical protein El 114503 Bloms anaises 34 78		153/3 1733 kroqubis homeobox protein 3				5777 Die Die Die Lieute Lieute de Li	Į.	5 146442 hypothetical protein DKF 2p434F2427.1 - (96.57% human)		54616 Exti3 exostoses (multiple)-like 3					RIKEN CDNA 2310032D 16 neme		chromosome 20 open readino frame 107: similar to neuronal thread brotein (Horne senienal 60 %	RIXEN CONA 5730493819 (62 07%, Muse miseribia)				Seriora		61895 Wdrig WD repeat domain 10		ŝ	Uxs1 UDP-glucuronate decarboxy/ase 1	Rpo2-4				Snrpd3	A43344 synaptic vesicle protein SV2 (58% rat)		Nxf1 nuclear ma export factor	Manta mannosidase alpha manta	KIAA0298 hypothetical protein (human)-ribosomal protein L27a-suppression of tumoripenicty 5		Slo6a4 serotonin transporter sert neurotransmitter: encoding serotin			Cst8 cvstatin 8 (cvstatin-related epididymal spermatopente)		Rab3a rab3a member ras oncodene family		PIND RYTHAGI 18 IA. to remissable	Tennoeur court or action of the court of the		AUCUA ALPUNIUM CASSENE ADDI-PRING AND INCIDIOS A BOOK / FINCE MOUNTED A BOOK /	
Locus	1602	17254	71846	1398			20074	74902	73332	18131	78783	7030	163/2				62740	04/10	20/615		54616					74182		76426	74478			RYBOR Cont	06070	9000	9199			67883	20022					75209		53319 Nxf1	17155 N	117224		15567 Slc6a4			13012 Cst8		19339 Rab3a		66617	200	toan About	76553	) ) )
Cluster Access	12C110_MM 2	2 AK020444	3 AK005954	3 NM 010157	3 NM 021475	3 NM 010580				3 NM 008716			3 NM 008393	3 BC004690	3 AK007368	3 1.25890	S NAM COLLEGE			3 AK005939	3 NM_018788	3 NM_025383	3 NM_023371	3 NM_023805	3 AK006268	3 AK009137	3 NM 007964		3 AK017115	3 NM 011168	3 BC006075	3 AK011256	3 AKO05560	00000000	3 Ar 2950/3	3 BC004674	3 NM_013483	3 AK005536	3 NM 011293	3 NM 025419	3 AKO 15036	2 200000	3 NM 026095	3 AK015921	3 NM_025829	3 NM_016813	3 NM_008548	3 AJ307670	3 BC003209	3 NM 010484	3 NM 008078	Ž		3 AK013874				_	NA COROOS	3 AK012967	

cluster analysis I lung cancer hypothalamus

Description Mus musculus gene for Cox 17p. complete cds	2 caspase 2 casp2	zinc finger protein of the cerebellum zict		5 phosphoprolein enriched in astrocytes 15		urate oxidase ec 1.7.3.3	RIKEN cDNA 1110033O09 gene			pre b-cell leukemia transcription factor 2 pbx2	engrated 2	gnrh-gap encoding gonadotropin-releasing hormone and gn-rh-associated peptide gap precursor		t cytochrome P450, tarmiy 2, subřamiry u, potypeptide	Moderner inward recilier massium channel the			RIKEN cDNA 4830415F15 gene		purinentic recentor (family A group 5)		RIKEN cDNA 2310035M22 [Mus musculus]		Mus musculus Y-box binding pratein (oxyR) mRNA	ubquiin C	cholesterol 25-hydroxylase	phosphalidylinositol glycan, class m, ref:NP_077058.1 • PIG-M mRNA for mamosyltransferase [Rattus norvegicus] 96 %					reflige myteir prid hinding argein 2			sofute carrier family 16 (monocarboxylis acid transporters), member 7	lysosomal alpha-glucosidase			q300 q300 protein aa 1-77	Similar to G-protein-coupled receptor induced protein (40% Homo sapiens)	DAKA Kindian media menangan	ANA Unitarily Mout Police to a	Production and the STDEM general production of the state	information protein rich of the supplier of th	solute carrier (amily 30 (zinc transporter), member 6	insulin-tike growth factor binding protein-1	transcriptional intermediary factor beta (if1b; tif1 protein	vesicle associated membrane protein cellubravin vamplsynaptobrevin homolog
8	12366 Casp2	22771 Zlc1	_	18611 Pea15		22262 Uox	5			5 Pbx2	9 En2	14714 Gnrh		risia cypzur	18513 Kcni10			•		67 168 P2v5	•				3	Ch25h	mbid					25064 Coho2	18013 Neurod2		20539 Stc7a5	Gaa			Hpvc2		9				Sic30a6	lgfbp1	Trin 28	/amb3
Locus	123(			1981		2226	68695			18515	13799	1471	7000	101	1651			73862		67 168		66949	80744	unknown	22190	12642	67556					75064	18013	67236	20539	14387			15456 Hpvc2	74178	9240C Dhan	10100		3	210148 SIc30a6		21849 7	22319 Vamp3
Chuster Access 3 ABN47323 3 NM 008916	3 NM_007610 3 NM_011483	3 NM_009573	3 NM_020585	3 AK010939	3 AK008493	3 NM_009474	3 AK004078	3 AK004627	3 NM_008834	3 NM_017463	3 NM_010134	3 M14872	3 022015	3 NM 007788	3 NM 020269	3 NM_025989	3 AK021330		3 AF322375	3 AK011967	3 AK012969	3 NM_025863	3 NM_030560	3 L35549	3 NM 019638	3 NM 009890	3 AK018560	3 11/93	3 NW 010008	3 MM 015740	3 NM 013881	3 AK015789	3 NM 010895	3 AK004355	3 NM 011404		3 BC005782	3 NM 026040	3 NM 008283		3 AKUZUU33	NM OFFE	3 NM 025387	3 NM 009469	3 BC005753	3 NM_008341	3 NM_011588	3 NM_009498

Christ Access		į	Percentaliza
	24132 2	4132 Zfp118	Zinc finger protein 118
4 M19413			
4 AK003581	17191 Mbd2	Mbd2	methyl-cpg binding protein mbd2 contains repetitive sequence derivert from cpg Island Itanslation
4 NM_025540	66402 Sh	£	sarcolipin
4 NM 010708	16859 1	Lgals9	lectiv, asiariose birdina, soluble 9
4 NM 011598	21884 F	edde-	perforalorial protein perf 15 lipid binding homolog cytoskeletal component with possible membrane role; riken cdna 1700007p10 1700007p10;k
4 NM_014193	12292 Cacnats	Secrets	edidim channel, voltaged-dependent. L'type, alpha 15 subunit Cristi unitata Cast sometis au
4 NM 021312	57750 W	Vdr12	wd repeat domain 12 done mgc 1938; nuclear protein vfm1p
4 BC004085 4 AK010336			
4 NM_011623	21973 Top2a	op2a	topolsomerase dna ii alpha top.2a
4 NM 030248			
4 AKD15181	14688 Gnb1	Į.	guanine nucleotide binding protein beta gmb f
4 NM 011141	18991 Pou311	ou311	pou domain class 3 transcription factor pou3f1; cct-6 octamer bindina protein
4 AK016149			
4 NM_011990	26570 51	lc7a11	solute carrier family 7 cationic amino acid transporter y+ system member slc7a11
4 AK005146	76499 CI	lasp2	CLIP associating protein 2
4 AK012868	10/581 Col18a1	0,1891	11 days embyo wien cdna chone: 2700007112 Delicen John 24 thorus in many
4 NM 019410	18645 Pf	'n	
4 NM 009501	22326 Va	- X	ventral anterior homeobox-contabing gene 1
4 NM 007817	13107 Cyp2f2	rp212	cytochrome p-450 naphthalene hydroxylase
	67315 Ceacam12	sacam12	CEA-related cell adhesion molecule 12-RIKEN cDNA 1600031J20; Ceacam12-C1; Ceacam12-C3 (Mus musculus) 100 %
5 BC008225 5 AK001742			
	75870 Tcamt	) EE	testicular cell adhesion molecule 1
5 NM 018873			
5 NM_011917	24128 Xn	Xrn2	5-3 exoriborardease 2
5 AB041350	12830 Co	Col4a5	col435 type iv collagen alpha-chain
5 NM_013597	17258 Me	Mef2a	myocyte entancer factor za metza
5 NM 010919	18088 NIC	Nkx2-2	nk2 transcription factor related locus 2 drosophila nkx2-2
5 KM 011031		P4ha2	procollagen-proline 2-oxoguidaate 4-dioxygenase proline 4-hydroxylase sipha ii polypeptide p4ha2
5 AK016/18	71062	1.10	KIKEN CUNA 483407/G/J gene
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5 NM 018749	٠ ٦		ing groces of process of the second sector of sectors o
5 BC003908			Council of the Counci
5 AK020790	77782 Polg		ONA directed), theta
S NM_019753	12557 Cdh17		CADH, MOUSE Cadherin-17 precursor (Liver-Intestine-cadherin) (L1-cadherin) (BILL-cadherin) (P130) 100 %
5 NM 007946	13861 Epx		eosinophil peraxidase
5 NM_030880			protein kinase C and casein kinase substrate in neurons 3
5 NM 009621	11504 Ada	Adamis1 8	incomplete and metalloprofease reprofysin type with thrombospondin motif adams 1; secretory protein containing motifs putative
5 NM 026089	67281 Buil37		stranger contained as a source of the contained and a source of th
5 NM 013867			
5 NM_010836	17703 Msx3		msx3 the drosophila melanogaster muscle segment homeobox msh protein encoded by genbank accession number u33319
5 NM_007667	12564 Cdh8		саднет-в
5 NM_012045 5 NM_015809			
5 NM 009575	22773 Zic3		zic3 pratein
5 AK002465	, 1001	•	The fact of the state of the st
5 AK007298	76999	. ¥	rnicen cours, sobustatics game KKK9 MOUSE (slandular kalikieni K9 precursor (Tissue kalikielin) (MGK-9) (Ebidermal growth factor-bi 68 %
5 NM_031873	83770 Tas1r2		candidate taste receptor 11/2 g protein coupled

cus Gene Description 229487 Pet 1121 PET 112-like (yeast) 75039 RIKEN cDNA 4930505H01 gene 74901 RIKEN cDNA 4930455M17 nene	Pcdh12		76730 RIKEN cDNA 2310005C01 gene 56358 Moret 7 adult mala lovena ditan cdoa ctora ctora 2310032m0			54427 Dnmt31 DNA (cytosine-5-)-methytransferase 3-tike		Similar to HUMAN Stress-Induced-phosphoprotein 1 (ST11) (Hsp?0/Hsp90-organizing protein) [44% Human] stip!		191 RIKEN CDNA 4921517J08 gene	53858 ORF5 open reading frame 5	Lcn4	14674 Gna13 guanine nucleotide binding protein alpha 13 gna13	56626 Poli polymerase (DNA directed), lambda	02 protein DKF2o56611024.1 - human (fragment) 137% human)		27 hypothetical protein, MNCb-2622	2584740 WK7171-12 domesticast done organ altadore deceptor, odorant k12 7 1660 Ferra 1 septimentals bitmen titlen entre chose includible altadore deceptor.			Mmp11	AIKEN CDNA 280001412	Alines		Itgam	Sp1	Smc411	×660		)	7 Mid2 midline 2	Hbs11		D C	6 Knil 4 10 days ambyo niken cida done-250060453; irad type i arasisk Mas muscrins adult mala testis chiva Dirkt N intanchi amadea iban dan attanchi amadea ilah	_	_		789.20 Inter order a Baloo-Zeboko done Inggr: (933) 1793.7 Tiona h media-interaction median do tindo	3	
Locus 229487 75039 74901	2360					5442		69545		70891	5385	16821	1467	<b>2662</b>	67902	70090	58227	71690	27214		17385	67107	102160	70315	16409	20683	70099	18292	17751		23947	56422	75396	74091	96736	70719	75727		64723	68501	
Cluster Access 5 BC005709 5 AK015703 5 AK015509	5 AK011541 5 NM_017378	5 NM_010218 5 AK010396	5 AK009163 5 NM 025450	S NM_009378	5 AK014062 5 AK015396	5 AJ404467	5 NM_019535 5 NM_009851	5 AK009368	5 AK005680	5 AK011489	5 NM_016924	5 NM_010695	5 NM_010303 5 AK012521	5 NM_020032	5 AK013017	5 AK009168		5 AF 282272 5 NM 023612		5 AK019926	5 NM_008606	5 NM_025971			5 NM_008401	S NM_013672	5 BC005507	5 NM_008759	5 NM 013603	5 AB044560	5 NM_011845	5 NM 019702	5 NM_029269		5 AK006831			5 NM_019519		5 AK003685	

Description		spectrin bata 4	DVS27-related protein [Homo sapiens] 51 %	dual adaptor for phosphotyrosine and 3-phospholnositides 1			ein General and	or the or partial transfer of the or the order of the ord	ULIANDALI	RIKEN CDNA 4921509B22 pene	RIKEN CONA 497517 INS none			hras suppressor expressed in skeletal muscle heart brain and bone marrow; 18 days embnyo diken edna done: 1190010123	forkhead transcription factor forest forest files				4 ribosomal protein S6 kinase	orthoradia homoton descendita ote	din sandara Area and area	nydroxy-deita-5-steroid denydrogenase, 3 beta- and steroid deita-isomerase 7	B lymphocyte activator macrophage expressed: BCM-like membrane protein (75% hyman)	hydrybairal malain MCC16733 (89%, human)			fring [Homo saplens] 83 %					54393 Gabbr1-Ubc gamma-aminobutyric acid (GABA-B) receptor, 1-ubiquitin D	Similar to hundipatival another El 19 (72) 1079, House enclosed	Common of placement process and the state of	OMININA GECADOXYASE, STACHLIAI	adult male testis riken odna clone:1700023a16		soling descention along training at a second of the second	Termin defected and the service of t	membrane-associated tyrosine-and threonine-specific cdc2-inhibitory kinase	secretory protein bv8	colicase parcreatic	hormonically interested May, consociated blaces	יייי איל אור איני האיני איני איני איני איני איני אינ	receptor tyrosine kinase-tike orphan for 1 - KOK1_MOUSE_1yrosine-protein kinase transmembrane receptor ROR1 precursor (Neurotrophic tyrosine kina 100 %	CASP2 and RIPK1 domain containing adaptor with death domain	mitochondrial ribosomal pratein L51	odd Oziten-m hamoloo 1 (Drosophila)		iliny ada desaturasa 1; deria-5 rany ada desaturase; imokeyy-CoA desaturase (delia-6-desaturase)			147169 hypothetical protein DKFZp762D096.1 - human (fragment) 82 %	molecule possessing ankwin-repeats induced by lipopolysacchanide mail	Mus musculus adult male diencephalen cDNA. RixEn full-length enriched library chose 93301611 no envelocitional accession	profess related with prophess (450k human)	יות משנים אותי של אותי	programmed ceu dean z pacaz	RIKEN CDNA 5530600A18 gene	origin recognition complex subunit orce		actions along	activities after a	hiproblast growth factor 17	paired related homeobox 1	ribonuclease 6 precursor (Homo sapiens) 67 %	not the manifest Harm contact 18 %	the remaining from softens for a	Holocytochome c synthetese	
Locus Gene		80297 Spnb4	77125	26377 Dapp1	28146 D3Ucla1	26443 Psma6	Stack Conc	10024 Day	BOY PCVIII	70859	70891	00000	87580	27281 Hrasrs	30923 Foxe3				56613 Rps6ka4	18420 Oro	700000	101302 HS030/	74748	101861	20000	27-114 6 1901	67338					54393 Gabbri-	70088	40000	18783 Odc	69371		40123 044	200	268930 Pkmy11	50501 Prok2	109791 Clos	26550 Hunk	20653 0011	102 50007	12905 Cradd	66493 Mrpl51	23963 Odz1	70007	10701			72244	80859 Mail	77225	69401		2000 F0002	71452	56452 Orc6		11474 Acts	2007	14171 rgm/	18933 Pro:1	68195	66684	20000	15159 HCCs	
Cluster Access	5 AK008318		5 BC003847	5 NM 011832	5 NM 030685	5 NM 011968	5 NM 000119	S NA4 000038		5 AK014850	5 AK014910	S AKONSESO	OCOCOON F	5 NM 013751	5 NM 015758	5 AK017667	100000	S NM DCDDC	5 NM_019924	5 NM 011021	£ 0€377740	9 77 77 10	5 AK017911	5 BC013813	9 104 040000	20000	6 NM_026097	6 NM 019933	6 AKO12087	10071070	D ANDZOS//	6 AL078630	6 AK009177	C 404 042044	6 NM 013014	6 AK006257	6 109638	A NAM ACREGA	DOSOON MINIO	6 NM 023058	6 NM_015768	6 NM 025469	A NM 019755	S NIN 013046	CASC O MINI O	068800 WN 9		6 NM 011855	8 AVOCOOTE	0 2000000	B AKOU9332	6 AFZ1/002	6 BC010478	6 NM_030612	6 AK020378	A AKODROO?	2000 184 9	0 MM 000/33	6 AK017451	6 NM_019716	6 AK016041	A NM OLIVER	700000	DOODO WAY	6 NM_011127	6 AK015947	6 NM 025203	COLCADO MINIO	DELEGIST OF	

ene Description adult male lestis riken cdna clone:1700123m18	ex 15 testis expressed gene 15; 2210014E14Rik (Mus musculus) 100 %			sa4 serum amyloki a saa4 pseudogeneand saa5 pps3 COP9 (constitutive photomorphogenic) homolog, subunit 3 (Arabidopsis thatiana)	in 1 strongl interaction molecule stim 1	intergral membrane protein km t	ppp aspany amhopepidase am Er neemon InA InM bish affinio			171 dert i stake des monove ecceptor tras condante gette mouis Mar natscules adult male himocamens CDNA RIKEN full-kendet endriched fürzer, chens 2900024918 renduct beweites Denfilmbilenen contain mentein	AD21 protein (88% human)		1920 matcophage infamentatory proteint 3 alpha cor chemokine mill-3; land management of the match protein internation of the second and second and second and second and second s			hypothetical protein FL/13283(71% human)	sfi			44 milativo n-mulain resunted recentled recentled recentled recentled recentled recentled recentled recentled				tt msmxt sording nexin; snxt			11 TAF11 RNA polymerase II, TATA box binding protein (TBP)-associated factor		к даптпа-фитапул сатьохујаѕв		e3 forkhead box e3 foxe3			12 chemokine (C-X-C molit) ligand 12, (81182 cytokine - mouse 100 %
Locus Ge Unknown	104271 Tex15	17025 Alad 17087 Ly96	66349 13618 Ednrb	20211 Saa4 26572 Cops3	20866 Sti	16430 Ilm1	13437 Dnpep		21385 Tbx	13/33 Emr1 72871	68659	12532 Cdc	20297 Scya20			66039	Ē			14764 Gnr	16180 Illrap			56440 Snx1			68776 Taf1		56316 Ggcr		15377 Foxa3			20315 Cxd1
Chater Access 7 AK007258 7 NM 013501	7 NM_031374 7 AK018179 7 NM_053266 7 BC002251 7 AB035381	7 NM_008525 7 NM_016923	7 NM_007904	7 NM_011316 7 NM_011991	7 NM_009287 7 AK008402 7 NM_013804	7 NM_008408	7 NM_016878 7 AB048834	7 NM 013782	7 NM_009324	7 AK013595	7 AF285091	7 NM_009860	7 NM_016960 7 NM_007733	7 AK005253	7 NM_013470	7 AK002441	7 NM_031999	7 AK016603 7 AB047820	7 NM_019983	7 NM 009962	7 NM_008364	7 AK005049	7 NM_025400	7 NM 019727	7 NM_026401 7 V00711	7 AK014639	7 BC005603	7 NM_007440 7 X76011	7 NM_019802 7 BC019215	7 NM_011202	7 NM_008260 7 NM_008593	8 NM_008233	8 NM_008136	8 NM_021704

cluster analysis I lung cancer hypothalamus

Locus Genre Description	16351 1pp actin-binding protein mipp	20908 Stx3 syntaxin 39		66695 Aspn asporin precursor aspn preproprolein type i extracellular matrix leucine-rich repeat protein 53975 Dox20 DEADAH (Asp-Glu-Ala-Asof-His) box polyneotide 20	Cod7			SS214 Scamp4 secretory carrier membrane protein 4		1/2556 Cd716 KSp-Caddelin Comits and Comit Comits and C			Pik3ca	0.029 FINEST CATANTA SET LOADANT 1988 AND STARS AND STARS AND STARS AND SET LOAD AND SET LOAD	Golosia		Gpr34		1252 I Kai i c33/r2/ta4 homologue of the c33 antigen a target mabsimibilory to httv-l induced syncitum formation	60178 Cours enditor newin s	Atn8b2		18777 Lypia1 hysophosipase 1			80877 Lrb3 (ab3du protein	3818 Edn3 Edn3	_		88828 Sync syncolin dystrobrevin binding protein		Trim34	Mapk3	Tecot		77749 hypothetical protein FLJ 1254 (38% human)		11.564 Action medium-cova deliyodigelase accion			66091 Ndula3 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3	8137 Bahata - rahita member of ras menopra family		
Access	9 AF285178	9 D29797	9 NM_026111	9 AF316825 9 NM 017397	9 AF080580	9 NM_008458	9 NM 009309			9 NM 007663	9 AK005610	NM_023526	NM_008839	B ARUTTA	NM 013747	AK009293	NM_011823	9 AK005989	9 NM_007656	NM 024225	AF156546	NM_023135	NM_008866			9 NM_030695 8		9 NM 011979		NM_023485	NM 010174	NM_030684	BC013754	NM_021053	NM_020588	AK020339	BC009123	NM 00/382	AKU1326/ AF285577		AK006243	NM 011228	AK019981	NM_029121
Chuster															_		•		<b>u</b>			Ψ.	J.	<i></i> (	(	⊸ oπ		, ct	6	6	6	Φ.	<b>О</b>	6	<b>O</b> D 1	Φ.	<b>.</b>	<b>*</b>	P 4	n on	2 9	2 2	2 9	2

cass Locus Gene Description (007484	M, 011883 24017 Rnf13 ring finger protein 13 M, 054097 117150 pip5k2c phosphatidylinositol-4-phosphate 5-kinase. type II. gamma M, 013730 27218 Slam signatling fymphocyte activation molecule	World May 19929 And 1992 And 1	13078 Cyp1b1 12402 Cbi	14533 21898 Tirk tol-like receptor 4  023057 65964 Zak mitk alpha  011550 21428 Tcfl4 transcription factor like 4 tcfl4  0110510 13118 Cyp46 cytochrome P450, family 48, subfamily a, polypeptide 1  010010 14179 Fgl8 fibroblast growth factor 8	202574 26968 Isir immunoglobulin superfamily containing leucine-rich repeat 2025774 2009368 2026315 57272 Ora16 gene for odorant receptor A16 2026315 57272 Ora16 gene for odorant receptor A16 2026315 37272 Ora16 gene for odorant receptor A16 2026315 37272 Ora16 gene for odorant receptor A16 2026315 37272 Ora16 All Structure And Structure All Structure And Structure And Structure And Structure And Structure And A1700100422 gene 2026315 77520 RIKE A. DANA 4700100422 gene 2026316 77627 RIKE A. DANA 4331407K02 gene 2026316 77627 RIKE A. DANA 4931407K02 gene 2026317 77627 RIKE A. DANA 4931407K02 gene 2026317 77627 RIKE A. DANA 4931407K02 gene	11832 72465 Zfp131 zinc finger protein 131 11832 72465 Zfp131 zinc finger protein 131 11832 72465 Zfp131 zinc finger protein 131 118021 70954 adult male testis riken cdna ctone.4922502b01 1009550 22678 Zfp2 zinc finger protein 2 zfp2; msz/87 1009571 57344 Cyrl9 methyltrasterase Cyrl9 1359 E215 e21 transcription factor 5 clone mgc:6043; e21-5 protein 100940 75564 Rikel choloszylvi gene 1359 E215 e21 transcription factor 5 clone mgc:6043; e21-5 protein 13197 66390 CGI-107 protein (95% human) 13197 66390 CGI-107 protein (95% human) 13197 66390 zerebellar feucine rich acidic nuclear protein lanp phosphoprotein 32 mpp32 encoded by genbank accession number v734; anp32 11737 Anp32 cerebellar feucine rich acidic nuclear protein lanp phosphoprotein 32 mpp32 encoded by genbank accession number v734; anp32
Cluster Access	10 NM_011883 10 NM_054097 10 NM_009441		10 NM_C00994 10 AK008287 10 AK008287 10 NM_0019765 10 NM_019765 10 D50080 10 AK008389 10 AK002109 10 AM_019776	10 AK014533 10 NM_023057 10 NM_011550 10 UM_010010 10 U18746	10 NM_025940 10 NM_025774 10 NM_025774 10 NM_020515 10 NM_020515 10 AK015309 10 AK013044 10 AK016516 10 AK016516	10 AKO 16531 10 AKO 1832 10 D 13695 10 D 13695 10 AKO 6550 10 AKO 6551 10 AKO 6540 10 AKO 13197 10 AKO 13197 10 AKO 13197 10 NM_021099 10 NM_021099 10 NM_021099 10 NM_027099

Chaster Access 10 AK016214 10 AF398969 10 AB009392 10 NM_026423	Locus 15388	us Gene 15388 Hnrpl	Description protein I mpd: ribonucleoprotein
	27393	27393 Mrpl39	10 day old male pancreas riken cdna done:1810033d11; unknown c21orf8
10 NM_019642 10 NM_016973 10 NM_023284	20014 50935	20014 Rpn2 50935 Siat7f	ribophorin II sialytransierase 7 ((alpha-N-acetyfneurominyl 2,3-betagalactusy)-1,3}-N-acetyf galactosaminide alpha-2,6-sialytransierase) F
10 NM_010447 10 U62673	15382	15382 Hnrpa1	rearranged ma binding protein fi-2
10 NM_020514 10 AK019654	78911		T09013 RING fineer protein Fxv - 24% mouse
10 NM 025301	27397	Mrp117	ribosomal protein mitochondrial 126 rpm126
10 NM_021344	57816 Tesc	Tesc	lescalcin lesc
10 NM 029926	266632 Irak4	Irake	interleukin-1 receptor-associated kinase 4
10 AK021409			
40 MAK 024297	80400		14.c marco due huncitadiscal escribativa deben 2.55 (ABRITO168)
	54624		mus muss cuts i grounded protein, come a confidence of the confide
10 U79525	14747	Cmklr1	orbitan d-protein coupled receptor dez
10 NM_011585	21677	21677 Tead2	TEA domain family member 2
10 AK004618	74106		RIKEN CDNA 1200006M05 gene
10 BC009093	13654 Egr2	Egr2	early growth response 2 done mgc;7113; zinc finger protein krox-20 exon a
10 NM 009871			
10 AKD15327			
10 AKD18520	71544		GTPase regulator associated with the focal adhesion kinase pp.125 (Homo sapiens) 56.19 %
10 AK019605	78799		reliction 1; neuroendecrible-specific protein (GSF, human)
10 BC016434		Maß	t-censectation from matter states and control of the control of th
10 NM 008123		Gjaß	gap junction membrane channel protein alpha gjaß
10 AB042432			
10 NM_026405	67844		10 days embyo fiken ddna chone;2810011317
10 NM 011342	20333 Sec2.	20333 Sec2211	SECUL VESTED Englishing professional (1), cereavillas)
11 AKOOB338	200	7054	
11 NM 020501			
11 AK014581			
11 AK009834			
11 NM 011787	23802 Arnth	Armth	autocrine motility factor receptor clone image:3500628; amit
11 NM_011821	23888 Gpcs	9	glypican is gloce
11 L13204	15223	Foxi1	hnt-3/forkhead homolog 4 hfh-4 bp 256303 activation domain region il 508810 winged helix dna binding
11 NM_011602	21894	拒	latin
11 AB016602			
11 AK014667	74577		Typerference protein McL 10 V / 1 (8 % Monto Sapiers)
11 NM 008166	14803 Grid1	Grid 1	adua inisier essis inisier odi accioni della contra
11 AK010153		;	
11 NM_010210		Ŧ	fragile histidine triad gene
11 NM_016972	50934	Slc7a8	gycoprotein-associated amino acid transporter laf2
11 AK013511		Ndut/2	NADH dehydrogenase (ubiquinone) flavoprotein 2
11 NM_007870		Dnase 113	drase gamma deoxyritonuclease
878520_MN 11	0/1/0	;	After EUNA 2 (VOU DECU GUERE dand shows however of Processes is a
11 NM 020483	57230 Mcnon	4000	11 dave embroo riken odne 270016405 fill incert semienrer hroni 11 dave embroo riken odne 270016405 fill incert semienrer hroni
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Similar to ENP1, HUMAN Ectonucleoside triphosphate diphosphotydrolase 1 (NTPDase1) (Ecto-ATP diphosphotydrolase) 44% uncharacterized bone marrow protein BM033 [65.88% Homo sapiens]
                                                                                                                                                          ovalburnin upstream promoter transcription factor i coup-tif i submitter comments: homodimer subunit s; coup-tit transcription factor 7-like 2, T-cell specific, HMG-box inorganic pyrophosphalase [Homo sapiens] 71 % RIKEN cDNA 2410157M17 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            signal sequence receptor, beta serineAhreoning (STK3; protein kinase homolog [Mus muscutus] 100 % ankyrin repeat, family A (RFXANK-like), 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                trans-premyfransferase
strain cut. I k-dependent na/ca exchanger stc24a3 ncbu3; sodium calcium clone image:3599824
RIKEN CDNA 2310010M24 gene
chromosome 20 open reading frame 108 [Horno sapiers] 77 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          intranuclear protein
protein phosphatase 1g formerly 2c magnestum-dependent gamma isoform clone mgc:6666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similar to putative breast adenocarcinoma marker (32kD) [Homo sapiens] 99 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               retinitis pigmentosa 9 homolog (human) transcriptional co-activator with PDZ-binding motif (TAZ) [91% Homo sapiens]
                                                                                                                                                                                                                                                                                                                                                                                                                      hoxb9 hox-2.5 abdominal-b homotog; protein cancerñeslis antigen 1; cancerñestis antigen 1; cancerñestis antigen [Homo sapiens] 36 %
                                                                                                                                                                                                                                                                                                       antigen identified by monoclonal antibody MRC OX-2 recepto
                                       17152 Mak male germ cell-associated kinase
54393 Gabbr1-Ubc gamma-aminobutyric acid (GABA-B) receptor, 1-ubiquitin D
22333 Vdact voltage-dependent anton channel vdact
                                                                                                                                                                                                                                                                                                                                                                           lysosomal-associated protein transmembrane 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            phosphaticiic acid phosphatase 2a ppap2a
nesclent helix toop helix 1
F-box only protein 12
                                                                                                                                                                                                                                                                                                                                                  poly(A)-binding protein, cytoplasmic 4-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chemokine (C-C) receptor 3
                     scrapie responsive gene 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     preprosomatostatin
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56075 Tptf
94249 Sk24a3
71897
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30924 Angpti3
19012 Ppap2a
18071 Nhh1
50757 Fbxo12
                                                                                                                                                             13865 Nr2f1
21416 Tcf712
                                                                                                                                                                                                                                                                                                                                                                                16792 Laptm5
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70544
20604 Smst
16328 Inmp
14208 Ppm1g
    20284 Scrg1
                                                                                                                                                                                                                                                                                  17527 Mpv17
                                                                                                                                                                                                                                                                                                                                                                                                                                               66192
12771 Ccr3
                                                                                                                                                                                                                                                                                                          57781 Mox2r
                                                                                                                                                                                                                                                                                                                                                                                                                        15417 Hoxb9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17922 Myo7b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            66256 Ssr2
56274 Sik3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   55934 Rp9h
                                                                                                                                                                                                            74776
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                                                                                                                                   11 NM_011720

11 NM_011720

11 NM_0209333

11 BC011417

11 AK010821

11 NM_020048

11 NM_0208622

11 NM_021325

11 AK013916

11 NM_024231

11 NM_024231

11 NM_024231

11 NM_025448

11 NM_025448

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11 NM_018501

11 NM_018501

11 NM_018503

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11 NM_018503

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11 NM_018503
11 NM_009136
                                            11 NM_008547
11 AL078630
11 NM_011694
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11 AK0087267
11 NM_026526
11 NM_025394
11 NM_025394
11 AK0002826
11 NM_01686
11 NM_01686
11 NM_01686
11 NM_01686
11 NM_01680
11 NM_033146
11 AK002327
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11 AK007580
11 AK006525
11 AK017626
11 NM_009215
11 NM_009215
11 NM_00814
11 NM_006106
11 NM_010616
11 NM_010626
11 NM_010626
11 NM_010626
                                                                                                                   11 NM_021336
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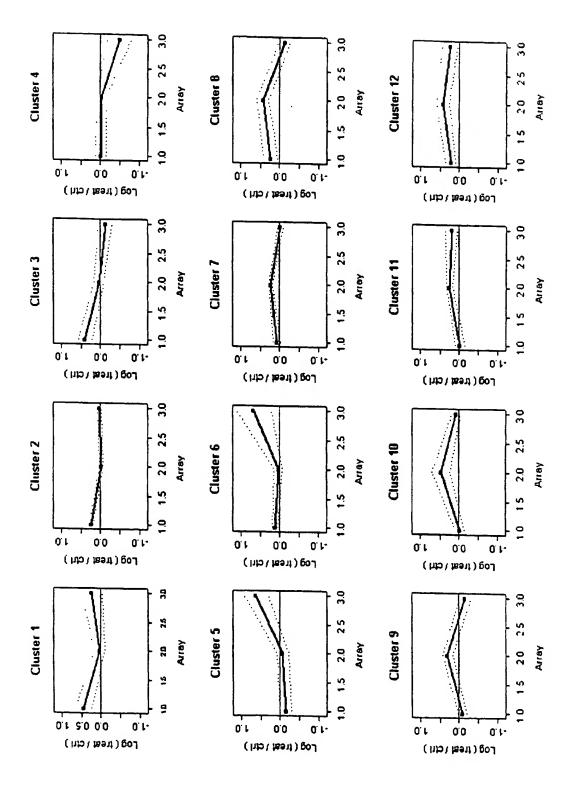
Description

Gene

Locus

Cluster Access

ionidination alian countries		0.0000 [			17765 MH2 metal response element binding transcription factor 2		1938)3 Sdnsl neural stem cell denved neuronal survival protein		21784 Tff1 treftoil factor1/5s2 exons 1-3	77422 RIKEN CDNA C330018D20 gene	27267 genomic fragment 281000 bp chromosome; cysteinyl-tma-synthetase cysrs	21855 Timm17b adult male tongue riken cdna done:2310005c24; 10 days embryo ckone:2610109e24	Wnt5a		19087 Prkar2a protein kinase, cAMP dependent regulatory, type II alpha		ECONO Comme at commercial annotational absorbance binding a fall to an over heat 100 M							71517 esophageal cancer associated protein (homo saprens) 92 %	19668 Ropsurt recombining binding protein suppressor of hartess-like (Drosophila)		68038 hypothetical protein MGC3234 [89% Homo saplens]			_	19250. Pipn 14 putative protein lyrosine phosphalase plp36 a sh3-binding site in the spacer region connecting n-terminal band 4,1-tike domain and c-terminal p	11888 Dahata - ashata mambaras famili dana meri 14577, as palis sitan dana dana dana da 14000 ana		58149 Grasp brain cdna done mrcb-4428 grp1-associated scaffold protein grasp unnamed product		75110 L1 repeat, Tf subfamily, member 29 [69% Mus musculus]			g .	14789 Gna1 giutamate receptor ionotropic empa1 alpha gna1		1570R Hash hairy and enthanner of entit & denomblis back	200		80907 Lacto serine beta lactamase-like protein lact-1		65591 Tmp21 Itananembrane Hafficking protein	7,2058 KIREN GUNA 20 IUU0,3D.20 gene
11 AK005633	12 AF359382	12 AK006008	12 AK020723	12 AK003179	12 NM_013827	12 AK017418	12 BC003996	12 AK010426	12 NM 009362		12 AJ276505	12 AF 106621		12 AK014861		12 BC018220		12 AKOOA33	12 NM 010253	12 AK013927	12 AK004009	12 AK015973	12 NM 023179	12 AK011615	12 NM 009036	12 AK005168	12 BC006876	12 NM_008292	12 NM_011041	12 X81466	12 NM 008976	081020 MN 21	12 AK008023	12 NM 019518	12 AK019591	12 AK015778	12 NM 019709	12 AK009010	<b>3</b>		12 AK003123		21 POLO TAIN CO	12 AK003278		12 NM_013582	12 AK003676	12 AKUUSUBU



Gene Description	_	hypothetical protein 1-82 [100% Mus musculus]	Parva parvin, atpha	Gjet gap junction membrane channel protein epsilon 1	hepatitis C virus core-binding protein 6 (70% human)	Similar to hypothetical protein FLJ11259 (Homo sapiens) 93 %	hypothetical protein FLJ22353 [Homo sapiens] 85 %		Fktp8 FK506 binding protein 8	_	T12515 hypothetical protein DKFZp434B103.1 - (28% human)	Pmftn1 polyamine modulated factor 1 binding protein 1				H 13 histogrammatibility 13		Times transfers of time mitochondrist membrane 9 homolog (veast)	_	polycythemia rubra vera 1; cell surface receptor; cell surface receptor [Homo sapiens] 49 %	RIKEN cDNA C330006D17 gene	Area amphirequlin	_		_	_	_	_	Nab1 Ngfi-A binding protein 1		Similar to myeloid/lymphoid or mixed-lineage leukemia 2 [41% Human]	_	Rgs9 regulator of g-protein signaling 9-2 isoform rgs9 rgs9-2 isoform striatal-enriched atternatively spliced product	_	hypothetical protein IMAGE3455200 [84% Homo sapiens]		Pripm prolactin-like protein M	_	_	-	_	Similar to sperm adhesion molecule 1 (PH-20 hyaluronidase, zona perucida pindurg) (45% Homo sapiens)	FIXEN CDNA 2310021M12 gene	_	_	-	CA.4a GPI-anchored metastasis-associated protein normolog	•		- 2012ai - P.O. dominair, Gassy, S. & Sasyucaning factor in the miseriliset (O) 9.	Tenzip thousand repeat ordinal declarate soming protein through the control of th	2.12200A-2301 Bette from setstim FINA RIKEN full-handt enriched library efton:CO30040A15 mediert-hynothetical Nicemeeskie setsti mele commissetstim FINA RIKEN full-handt enriched library efton:CO30040A15 mediert-hynothetical	INDS INDSANDS SOUTH THE CAPPAS STREAM. THE TANK THE TOTAL STREAM	Niva malaat transcription factor. Y alpha		S10151 transforming protein Ilm - 56% mouse (strain balbic)
Locus	75106	56279	57342	118446	67391	71712	68777	69533	14232	77785	70892	56523	59007	22350	88018	14950	12614	3005	109085	68891	77616	11839	101513	110639	16502	16425	74239	27224	17936	56463	66765	21206	18739	11937	27973	71989	56635	14829	17451	13110	19733	77042	71904	210376	11979				•			01077	70153	18044	68999	74085
Cluster Access		1 NM 019833	1 NM_020608	1 NM 080450	1 NM 026126	1 AK004552	1 AK004164	1 AK009086	1 NM 010223	1 AK020734	1 AK014905	1 NM 019938	1 NM 021504	1 MM 012703	* BC046023	1 NM 010073	1 MM OOGBBG	1 NM 013895	1 AK012767	1 AK004472	1 AK021182	2 NM 009704	2 AK012535	2 NM 026662	2 NM 008421	2 NM 010582	2 AK006472	2 NM_013736	2 NM_008667	2 NM_019776	2 AK016775	2 AK017114	2 NM_011268	2 NM_007504	2 AK002742	3 AK010322	3 NM_019991	3 NM_00B177	3 NM_020021	3 NM_010008	3 NM_009060	3 AK014599	3 AK009450	3 AF073881	3 NM 00751	3 NM 008497				3 NM_011136		3 AK01233	3 AAGE 130	3 AK004729	3 44005303	3 AK016812

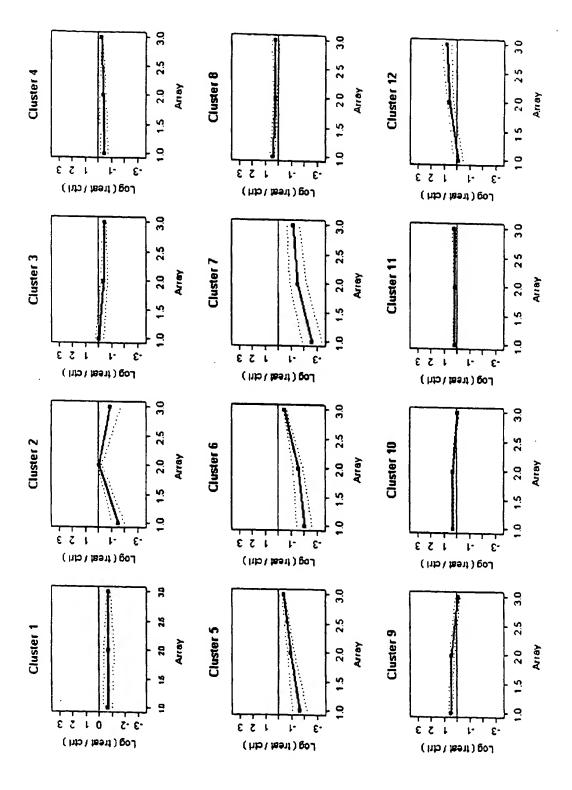
Gene	20916 Sucla2 succinate-Coenzyme A ligase, ADP-forming, beta subunit	Ahsg	Peti1	94228 Ucc1 upregulated in colorectal cancer gene 1	_	Lcn3	Pdo	V3R9		Drg2	_	Ş	Olfr159	Pgtyrpt	L EL	_		Bc110	Frap1	g		Bard	Caparapiz	1	12.0	MOVON	9	1947.4 Kgs to regulater of Cytotern signaling to		Psmc1		i	Zfp106	0493	QCIO PCIO		Sor3	Bnip3	N#DG1		Tor1b	14420 KIKEU CUNA 82540-VIG 88109 8810 8810 AND		VIRZ	17 FOUR MATILOS CHAINTONNAS C. AND MATILOS CHAINTONNAS C. AND MATILOS CHAINTONNAS C. AND MATILOS CHAINTONNAS C. AND MATILOS C.	0000	1 Inc 84a	
Chister Access	4 AF058955 Z(	4 NM_013465		4 NM_053192 84	_	394		2		4 NM 021354 13	4 NM 009491 22		9		4 NM_008707 18		4 NM_018832 54		Q.	4		4 NM_007525 12				o		_	AK007657		AK017569	AK017387	AF060248	NW 00//34	NM 009/44	AK009749	NM_009237	NM_009760	NM_010909	NM_026318	AJ297743		AK010800	-	2	5 AKG14490 (3130		

Chuster Access	Locus	9	Description
5 AK004654	71709		similar to A49307 98K GTPase-activating protein ABR, brain - human 29 %
5 NM 011374	20449	Siat8a	
5 AK012063	67181	Dullard	_
6 NM_025839	66911		
	66046	NdufbS	_
	14089	Fap	fibroblast activation protein
	66522		PGPI_HUMAN Probable pyrrolidone-carboxyfale peptidase (5-oxoprolyt-peptidase) (Pyrogiutarnyt-peptidas 95 %
	28998	<u>ਨ</u>	poliovirus receptor-related 3
6 AK005105	68980		Similar to WD domain, G-beta repeat-containing protein (Homo sapiens) 83 %
6 AF215896	11790	Apeg1	striated muscle-specific serineAhreonine protein kinase speg
6 NM 023175	52633		Nit protein 2 (89% Homo sapiens)
6 AK019095	78887		similar to pir:100322 - 100322 hypothetical protein KIAA0542 - human 62 %
6 AF056187	16001	lgf1r	insulin-like growth factor i receptor (gf-1
6 NM_009843	12477	CHad	cytatoxic T-tymphocyte-associated protein 4
6 NM_010351	14836	Ssc	goosecoid gsc
6 X58472	16588	ž	entigenic determinant of rec-A profein
6 NM_009763	12182	Bs:1	bone marrow stromal cell antigen 1
6 NM_023755	81879	C <sub>z</sub>	transcription repressor crti-1 developmentally regulated related to the cp2 family of factors
6 M12289	17885	Myh8	perinatal skeletal myosin heavy chain 3 end
6 AK011897	72495		RIKEN CDNA 2810208C17 gene
6 AF448804	72162	Ddx38	
6 NM_007657	12527	င္မီ	CD9 antigen
6 NM_053168	94091	Trim11	tripartite motif protein trim 1 t
6 NM_024465	76192		RIKEN CDNA 6330583M11 gene
6 NM_019435	104130	Np15	Nuclear neuronal protein 15.6
6 X72307	15234	ğ	hepatocyte growth factor
6 NM_080638	78388	Mvp	major vault protein
7 AK004076	68036		HSPC038 protein (Homo sapiens) 100 %
7 NM_026309	87678		LSM3_HUMAN U6 snRNA-associated Sm-like protein LSm3 (MDS017) 100 %
7 NM 026031	67205	Cgi94	CGI-94 protein
7 NM 027170	96969		JC6547 high sulfur protein B2E - rat 37 %
7 NM_011054	18575	Pde 1c	phosphodiesterase 1C
7 NM_022024	63986	Gmfg	glia maturation factor, gamma
7 NM 025816	52440	<b>36End772</b>	JGEnd772: Tax1 (human T-cell leukemia vinus type I) binding protein 1; tax1-binding protein (Horno sapiens) 80 %
7 NM 007507	11958	Atp5k	atp synthase h+ transporting mitochondrial f1f0 complex subunit e atp5k; tfm-1 f1f0-atpase
7 NM_019965	56709	Dnajb12	mdj10 deduced amino acid sequence homologous to c. elegans putative dnaj protein z73102 b0035.14. homolog
7 NM_016849	54131	E)-	interferon regulatory factor-3 inf3 inf-3; factor 3
7 AK020621	78670	Gurpx	guanina nucleotide releasing protein x
7 NM_023167	66163	Mrp14	mitochondrial ribosomal protein L4
7 NM_011870	23991	Cip	calcium and integrin binding 1 (calmyrin)
7 NM_008554	17173	Asct2	асhaete-scute complex homolog-like 2 (Drosoph#a)
7 AK007013	74282		
7 NM 030701	80885	Puma-9	pulative seven transmembrane spanning receptor puma-g
7 NM 028189	72297	B3gnf3	UDP-GicnAc: beta-1,3-N-acetyglucosaminytransterase 3
7 AK017362	75736	9	BCL2-like 12 (proling inch); Bd-2 like proline-nch protein 12 [Homo sapiens] 81 %
7 AK003011	17714	Grpeiz	Criterike Z, miochondria
7 NM 011658	22160	WSL	Wist gene nomotog 1 (Urosophila)
7 NM 025855	52665		unchairadetried hypothaamus protein HCLUASE [//% Homo Sapiens]
7 AK018541	1364		903000/LIVIK KIKEN CUMA BUJUGU/LIVI gene
7 AKUUDD / 0	321010	9	RINER CUNA. I GUILLE GUILLE
7 NM 021368	58168	9135	or order decelor of
7 NM 00/3/4	79771	5 E	Confidentent confidence of a Successformers galante polypetrule The Confidence of the Confidence of th
767600 MN 7	60121	CIGHO	grown diserentiation rectors so goi-so, one maybringered to only to
7 NM_009838	12466	8 5	Chaperon's supulin a (2013)
7 NIM 013483	1223	Birds at a f	Avenabilin subfamily I member A1
8 BC002307	27878		Outpring, account 1; married 7. SPT3-associated factor 42 [95% Homo sapiens]

hypothalamus

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Mouse 12 days embyo head cDNA, RIKEN full-tength enriched library, done:3010025E 17 product:(HNRNP A0) homolog [Homo sapiens]
Lysozyme C, type M precursor (1.4-beta-N-acetyfmuramidase C) (62% Mus musculus)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus adult male testis CDNA, RIKEN Mil-length enriched fibrary, clone:4930465412 product:hypothetical protein 
6-pyruvoyl-tetrahydropterin synthasedimentization collactor of hepatocyte nuclear factor 1 apha (TCF1) 
hypothetical protein MGC7537 [Mus musculus] 100 potassium large conductance calcium-activated channel, subfamily M, alpha member 1
                                                                                                                                                                                                                                                                                                                    ectonucleoside triphosphate diphosphohydrolase 5
Mus muscutus adult male testis cDNA, RIKEN full-length enriched library, done:4930448AZ0 product:unclassifiable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pir.114768 - 114768 hypothetical protein DKF2p566K1924.1 - human (fragment) 93.59 %
BCL2-like 12 (proline rich); Bcl-2 like proline-rich protein 12 [Homo saplens] 81 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          myelokanymphoid or mixed lineage-leukemia translocation to 3 homolog (Drosophila)
                                                                                                                                                                                                                                                                                                                                                                        Similar to seven transmembrane protein TM75F3 (82% Homo sapiens)
SNIL_HUMAN Probable serinenthraonine protein kinase SNF1LK (55% fruman)
hypothetical protein FLJ10560 (82.88Homo sapiens)
tor beta locus from bases 250554 to 501917 section 2 of 3 the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gamma-aminobutyric acid (GABA-A) receptor, subunit beta 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RAB1, member RAS oncogene family CU56_MOUSE Putative protein C21orf56 homolog 54.41 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          adult male medulla oblongata riken cdna clone:6330437i11
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (clone pcm7) growth hormone-releasing factor receptor
                                                                                                                                                                                                                                   ribosomal large subunit pseudouridine synthase C like
                                                                                                                                                                                                                                                                                                                                                            Y174_HUMAN Hypothetical protein KIAA0174 95 %
                                                                                                                                                                                                                     glycerol phosphate dehydrogenase 2, mitochondrial
                                                                                               RIKEN CDNA 4932442K08 [Mus musculus] 100 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical protein FLJ22724 [68% Homo sapiens]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        similar to S. cerevislae RER1 [Homo sapiens] 96 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PDZ domain containing 1Mus musculus) 100 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DnaJ (Hsp40) homolog, subfamily B, member 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KIAA0773 gene product [31.09% Homo saplens]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        heterogeneous nuclear ribonucleoprotein D-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuroblastoma, suppression of tumorigenicity 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lissue specific transplantation antigen P358
                                                                                                                                                                                                                                                                                 sushi-repeat protein (Homo saplens) 93 %
                                                                                                                   Ellis van Creveld gene homolog (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tissue inhibitor of metalloproteinases-3
                                                                                                                                                                                                                                                                                                   Cip1-Interacting zinc finger protein
                                                                           RIKEN cDNA 2010309E21 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       dystrophia myotonica linked gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   - I38487 tastin - human 51.90 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SYN1_MOUSE Synapsin I 30 %
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RIKEN cDNA 1200011118 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RIKEN CDNA 4632428N05 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A30411 synapsin la - 34% rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               interleukin 12 receptor, beta 1
                                                         oocyte specific homeobox 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            caudal type homeo box cdx4
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                                                                                                                                       testis-specific c-abl protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Msx-interacting-zinc finger
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               annexin A11
                                       contactin 2
                                                                                                                                                                                                                                                             cystatin 9
Gene
Dip3b
Cntn2
Obox1
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Pdzk1
Onajb8
Dm9
Tpo
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Gpd2
Csf3 bd2
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Cdx4
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70573
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AK010262
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AK06894
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1 NM_021517
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) AK018205
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                                       8 X81365
   Chuster Access
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Description	leucine zipper-EF-hand containing transmembrane protein 1	tocopherol (alpha) transfer protein	calpain 7	hypothetical protein FLJ22362 (98% human)	days neonate head riken cdna clone:5430425h06; 9130005n14 9130005n14 pix	signal sequence receptor, gamma (translocon-associated protein gamma) [Homo saplens] 98 %	done mgc:7273	signal sequence receptor, alpha	cytochrome P450, family 4, subfamily v, potypeptide 3	antigen p97 (melanoma associated) Identified by monoclonal antibodies 133,2 and 96.5	сулостоте Р450, family 2, subfamily d, polypeptide 26	hypothetical protein FLJ11101 [76% Homo sapiens]	caseinolytic protease, ATP-dependent, proteolytic subunit hornolog (E. coll)	ribosomal protein L36a; 60S ribosomal protein L44; L44-like ribosomal protein; ribosomal protein L4 (100% human)	OM07_HUMAN Probable mitochondrial import receptor subunit TOM7 homolog (Translocase of outer membran 94 %	CU59_HUMAN Protein C21orf59 92 %	180486 gene irg protein - 36% rat (fragment)	cd7 antigen	cDNA sequence BC005662	coagulation factor XIII, alpha subunit	RIKEN cDNA 1700025F22 gene
800	Letm1	Ttpa	Capn7				Sign	Ssr1	Cyp4v3	Mf2	Cyp2d28		ਨੂੰ					g		F13a	
Locus	56384	20500	12339	64339	68303	67437	83429	107513	102294	30060	76279	67509	53895	66483	69199	68001	67539	12516	210992	74145	69416
Chater Access	12 NM_019694	12 AK004882	12 NM_009796	12 NM 022424	12 NM 026667	12 AK002818	12 NM_031251	12 NM_025965	12 AB056457	12 NM_013900	12 NM_029562	12 AK015650	12 NM_017393	12 NM 025589	12 AK003864	12 AK003413	12 AK016777	12 NM 009854	12 BC/05662	12 NM_028784	12 AK006335



## Figure 24A

#### DIFFERENTIALLY EXPRESSED GENES (SELECTED)

#### Colon

Accesion	Gen	Description	Class	16
X72307	Hgf	hepatocyte growth factor		Region
U92885	Efna3	aphenn A3	growth factor	Cortex
NM_026908	Cts3	cathensin 3	growth factor	Cortex
NM_025896	plpi	prolactin like protein l	protein degradation (secreted)	Hypothatamus
NM 021704	Cxel12	chemokine (C-X-C moxif) ligand 12, 181182 cytokine - mouse 100 %	growth factor	Cortex
NM_021380	II20	interteukin 20 il 20	immune response (secreted)	Hypothalamus
NM_017383	Cntn6	contactin 6 - neural recognition molecule NB-3 (Mus musculus) 100 %	growth factor	Cortex
NM 016851	tr16	interferon reculators (notes a mide into moscories) 100 %	immuna response (secreted)	Midbrain
NM 013857	Sema3	interferon regulatory factor 6 mirf6 irf6 transcription; clone mgc:5918	immuna response (secreted)	Micorain
MM_013552	Cd4	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C chemokine (C-C motif) ligand 4	growth factor	Cortex
NM_011165	Prtpa	projectin-like protein A	growth factor	Striatum
0	lbe	protectiving protein A	growth factor	Midbrain
	j.			Conex
NM_010735	Lta	Landard to A		Striatum
MM_010/33	irea.	lymphotoxin A	growth factor	Striatum
NM 009971				Hypothalamus
NM_009971	Cs/3	colony stimulating factor 3 (granulocyte)	immuna response (secreted)	Cortex
WW_009121	8mp15	growth differentiation factor-9b gdf-9b; bone morphogenetic protein 15 bmp15	growth factor	Midbrain
			•	Cortex
NM_009640	Agpt	angiopoletin	growth factor	Midbrain
	1		grown ractor	
	1			Cortex
NM_009500	Vav2	vav2 oncogene	growth factor	Striatum
			growth factor	Midbrain
	ŀ			Contex
NM_009390	TO	tolloid-like		Striatum
NM_009129	Scg2	secretogranin (I	protein degradation (secreted)	Hypothalamus
NM_009043	Reg2	regenerating islet-derived 2	growth factor	Hypothalamus
NM 003930	Prine	prolactin-like protein E	growth factor (secreted)	Midbram
NM_008675	Nbit	neuroblastoma, suppression of tumorigenicity 1	growth factor	Cortex
	1	The second of th	tumor related (secreted)	Midbrain
NM_008382	Innbe	defense/immunity protein activity, indoleamine-pyrrote 2,3-dioxygenase activity		Hypothalamus
NM 008343	igfba3	insufin-like growth factor binding protein-3	growth factor	Striatum
NM 008108	Gdl3	growth differentiation factor 3	growth factor	Cortex
NM 008007	Fgf3	Fibrobiast growth factor 3	growth factor	Cortex
NM 008004	Fg117	libroblast growth factor 17	growth factor	Midbrain
NM_007995	Fcna	ficalin A	growth factor	Cortex
AK020723	I Cila		immune response (secreted)	Hypothalamus
AK020723 AK020305	I	2004395A melanin-concentrating hormone (100% Mus musculus)	growth factor	Hypothalamus
AK017955	l	sp.P11590 - MUP4_MOUSE Major unnary protein 4 precursor (MUP 4) 39 %	transport (secreted)	Cortex
WV011822	Myg1	melanocyte prolifeating gene 1	signaling (secreted)	Midbrain
	1			Cortex
	1		ł	Striatum
AK009012	Chit	chitmase (chitotrisosidase)	immune response (secreted)	Hypothalamus
AK008922	Fgf22	fibroblast growth factor 22	growth factor	Cortex
AF360358	ORF9	open reading frame 9	growth factor	Striatum
AF 158744	Pripc2	prolactin-like protein C 2	growth factor	Cortex Striatum

## Figure 24B

## DIFFERENTIALLY EXPRESSED GENES (SELECTED)

#### Breast.

Accesion	Gen	Description		
X76290	Shh		Class	Region
X72307	Hef	chymotrypsin activity, growth factor activity	growth factor	
X723111	rigi	hepatocyte growth factor	growth factor	Hypothalamus
	1		\$10mm ration	Midbrain
U92885	EIna3	apherin A3		Cortes
U56650	Nxph2	neurexophilin 2	growth factor	Cortex
NM 053192	Ucci	upregulated in colorectal cancer gene I	signating (secreted)	Michrain
•	1	ab eleganor in consecuti critical Bana i	turnor related (secreted)	Midbrain
NM_023517	Tofsf13			Conex
NM 023400		tumor necrous factor (ligand) superfamily, member 13	growth factor	Cortes
.4,41_02,54170	npc2	Narmann Pick type C2	miscellaneous (secreted)	
	1		imperiations (scenarion)	Midbrain
NM_021283	04	interleuxin 4		Hypothalamus
NM_020597	Msmb	beta-microseminoprotein; beta-inhibin; prostatic inhibin protein (Mus musculus)	growth factor	Midbrain
NM 019941	Pripm	protectin-like protein M	growth factor	Cortex
•	1	Promise process of	growth factor	Midbrain
NM 019775	Cob2	Company of the Compan		Hypothalamas
NM_019626	Chini	carboxypeptidase B2 (plasma)	protein degradation (secreted)	Midbrain
NM 017389		cerebellin 1 precursor protein	growth factor	
	Ear4	eosinophil-essociated ribonuclease 4		Midbrain
NM_017383	Critn6	contactin 6 - neural recognition molecule NB-3 [Mus musculus] 100 %	miscellaneous (secreted)	Hypothalamus
NM_015770	a	nonagouti	immune response (secreted)	Midbrain
NM_013652	Ccl4	chemokine (C-C motif) ligand 4	signaling (secreted)	Hypothalamus
• • • • • • • • • • • • • • • • • • • •		are meaning (C.C. made) algano e	growth factor	Cortex
NM_011888	Cd19	champling (O.C.) and the		Hypothalamus
NM_011187		chemokine (C-C motif) ligand 19	growth factor	Midbrain
	Pripc1	protectin-like protein C 1	growth factor	
NM_010504	Ifna4	interferon sipha family, gene 4		Cortex
NM_010203	Fgf5	fibroblast growth (actor-5 fgt-5 dna exon	growth factor	Midbrain
NM_010197	Fgfl	fibroblass growth factor 1	growth factor	Cortex
	1		growth factor	Midbrain
NM 009971	Cst3	colony stimulating faces a design		Contax
NM 000757	Bmp15	colony stimulating factor 3 (granutocyte)	immune response (secreted)	Cortex
TANDARY I	Cub12	growth differentiation factor-9b gdf-9b; bone morphogenetic protein 15 bmp 15	growth factor	
	1		Brown among	Midbrain
	1			Cortex
NM_009311	Tacl	tachykinin I		Hypothatamus
			growth factor	Midbrain
NM 009255	Scrpine2	estina (es parientes)		Cortex
	ac.p.i.c.	serine (or cysteine) proteinase inhibitor, clade E, member 2	protein degradation (secreted)	Midbrain
			, ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	Cortex
	1			
SM_009141	Cxd5	chemokine (C-X-C motif) ligand 5		Hypothalamus
NM_009042	Regi	regenerating islet-derived [	growth factor	Cortex
NM_008987	Ptx3	pentaxin related gene	growth factor (secreted)	Cortex
NM_008930	Pripe	profectin-like protein E	immune response (secreted)	Cortex
-	1	The state of the s	growth factor	Midbrein
NM_008734	Nrg3	and the second s		Hypothalamus
0007.04	nu go	neuregutin 3	growth factor	
			J. C.	Cortex
NM_008675	Nbit	neuroblestoma, suppression of tumorigenicity 1	Number and and of the con-	Hypothalamus
NM_008493	Lep	leptin	tumor related (secreted)	Cortex
			growth factor	Midbrain
NM 008343	lgfbo3	insulin-like growth factor binding protein-3	i	Cortex
NM 007423	Afp	alpha fetoprotein	growth factor	Midbrain
D01093	PCSk4		miscellaneous (secreted)	Midbrain
AK019795	resid	proprotein convertase subtilisin/kexin type 4	growth factor	Cortex
AKU14145	1	YA02_HUMAN HYPOTHETICAL PROTEIN DI 198H6.2 (50% HUMAN)	growth factor	
			Brown Istra	Midbrain
AK017955	Myg1	melanocyte prolifeating gene 1		Hypothalamus
AK015642	1	CERU MOUSE CERIA OPI ASMINI PRECUPCOR (CERRONIA CER	signaling (secreted)	Hypothatamus
AK013765	Ecgfl	CERU_MOUSE CERULOPLASMIN PRECURSOR (FERROXIDASE) (96% Mus musculus)	miscellaneous (secreted)	Cortes
AK009012	Chid	conduction cent growth factor I (platelet-denived)	growth factor	Midbrain
		chitinase 1 (chitotrisosidase)	immune response (secreted)	Midbrain
AK008922	Fgf22	fibroblast growth factor 22	growth factor	
	1		-	Cortex
AK006679	i	Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700041N15 product:	J	Hypothalamus
	1	minimum control interest in the control in the cont	Browth factor	Midbrain
AF210429	P1a2g10	phospholipase A2, group X		Cortex
		Thursday universe u.e. At AD V	miscellaneous (secreted)	Midbrain

## Figure 24C

## DIFFERENTIALLY EXPRESSED GENES (SELECTED)

## Lung

¥ 17107	Gen	Description	Cless	Region
12307	Hgf	hepatocyte growth factor	growth factor	Midbrain
	1		g.ova taztu	
	1			Contex
	1.			Stnatum
W2885	Efna3	aphenn A3	indust Comm	Hypothalamus
156650	Nxph2	neurexophitin 2	growth factor	Stratum
J18748	Fgf8	fibroblast growth factor 8	signating (secreted)	Striatum
			growth factor	Cortex
KM_053192	Ucel	upregulated in colorectal cancer gene i		Hypothalamu
-		The state of the s	tumor related (secreted)	Midbrain
	l .			Corner
NM_029269	1	Secreted abountains 24 programs (Con 24) (Con-		Hypothalamus
NM 028784	F13a	Secreted phosphoprotein 24 precursor (Spp-24) (Secreted phosphoprotein 2) (68% human) congulation factor XIII, alpha subusit	miscellaneous (secreted)	Hypothalamu
MM_026087	Ceacam 12	CEA related sell extreme makes to de Covers and a covers	immune response (secreted)	Hypothalamus
SM_023517	ToBO3	CEA-related cell adhesion molecule 12 -RIKEN cDNA 1500031J20; Ceacam12-C1; Ceacam12	tumor related (secreted)	Hypothalamu
05>>>>	111111111111111111111111111111111111111	tumor necrosis factor (ligand) superfamily, member 13	growth factor	Midbrain
	1			Cortes
VM_021782				Striatura
	1121	interlaukin 21	growth factor	Striatum
VM_021704	Cxd12	chemokine (C-X-C motif) ligand 12, I81182 cytokine - mause 100 %	immune response (secreted)	
	i		rese (estrate)	Cortex
M_021489	F12	congulation factor XII (Hageman factor)		Hypothalamu
VIM_021380	1120	interloukin 20 il20	misocilaneous (secreted)	Midbrain
	1		growth factor	Midbrain
VM_021283	114	interleukin 4		Hypothalamu
VM_021274	Cxd10	chemokine (C-X-C motif) ligand 10	growth factor	Midbrain
VM_020597	Msmb	beta-microseminoprotein; beta-inhibin; prostatic inhibin protein [Mus musculus]	growth factor	Midbrain
		A series in a series in a series by cores a light a turn a countries	growth factor	Cortex
VM_020021	Mos	Moloney sarcoma oncogene		Hypothalamu
	1	· · · · · · · · · · · · · · · · · · ·	growth factor	Midbrain
	l .			Cortex
VM_014991	Prtpm	prolactin-like protein M		Hypothalamus
	1	Protesting direct blocking Wi	growth factor	Striatum
VM_019450	11168	interded this distance is not a		Hypothalamus
NM_015770		interteukin 1 family, member 6	growth factor	Midbrain
NM_013657	Scma3	nonagouti	signaling (secreted)	Hypothalamu
M 013652		suma domain, immunoglobulin domain (Ig), short basic domain, accreted, (semaphorin) 3C	growth factor	Cortex
430 M 3032	Cc14	chemokine (C-C mouf) ligand 4	growth factor	Cortex
*** 01744	i			Stnatum
VM_013465	Ahsg	alpha-2-HS-glycoprotein	protein degradation (accreted)	Hypothalamus
NM_011697	Vegib	vascular endothelial growth factor B	growth factor	Michrain
NM_011423	Smr3	submaxillary gland androgen regulated protein smr3; msg1	growth factor	
NM_011330	Cd11	small chemokine (C-C motif) ligand 11	growth factor	Cortex
NM_011165	Pripa	prolacum like protein A		Micorain
			growth factor	Midbrain
NM_010253	Gal	galanin		Contex
NM_009757	Bmp15	growth differentiation factor-96 gdf-96: bone morphogenetic protein 15 hmp15	growth factor	Hypothatamu
-		15 pmp15	growth factor	Midbrain
	}			Striatum
NM_009732	AVP	arginine vasopressin		Hypothalamus
	1	manua seroherru	growth factor	Midbrain
NM_009704	Areg	amphires. La		Striatum
NM_009500	Vav2	amphiregulm	growth factor	Hypothalamus
	V8V2	vav2 oncogene	growth factor	Midbrain
NM_009362	Ters			Hypothalamu
		trefoil factor1/ps2 expns 1-3	growth factor	Hypothalamu
NM_009311	Tac1	tachykinin 1	growth factor	Cortex
NM_009285	Stc	stanniocaton	growth factor	Midbrain
NM_009255	Scrpine2	scrine (or cysteine) proteinase mhibitor, clade E. member 2		
ZW 004111	Cvels	chumokine (C-X-C motif) ligand 5	protein degradation (secreted) growth factor	Striatura
NM_009129	Scg2	secretogranin ti		Midbram
NM_COKA75	NHI	neuroblastoma, suppression of tumorigeniesty 1	growth factor	Cortex
	ì	The same of the sa	tumor related (secreted)	Midbrain
	1			Strietum
NM_008520	Lttp3	tatent transforming growth factor beta binding protein 3		Hypothalamus
NM (88447	Lhb	terrunting hormone bera brinding protein 3	signaling (secreted)	Striatum
VM_000351	11122	interfeukin 12 p33 subunu	growth factor (secreted)	Hypothalamus
			growth factor	Midbrain
NM_008004	Fgf17	fibrobless growth factor 17		Striatum
		Parametry of the Control of the Cont	growth factor	Midbrain
VM 007995	Fena	Seedin A		Hypothalam
	1_	ficolin A	immune response (secreted)	Midbrain
M148/2 D01093	Gnrh	gruth-gap encoding gonadotropin-releasing hormone and gruth-associated peptide gap precursor	growth factor	Hypothalam
	Pcsk4		growth factor	Striatum
AK020723	1	2004395A melanin-concentrating hormone (100% Mus musculus)	growth factor	
	1		Brown tarin	Midbrain
	l			Conex
AK017955	Myg1	melanocyte prolifeating gene 1	lementine tenner ::	Hypothalam
			signating (secreted)	Striatum
AK015842	Į.	CERU_MOUSE CERULOPLASMIN PRECURSOR (FERROXIDASE) (96% Mus musculus)		Hypothalami
	1	PERCONDUCT PROGRAM PRECURSOR (PERROXIDASE) (96% Mus musculus)	miscellaneous (secreted)	Midbram
	1			Contex
AK008922	Fg122	fibroblast growth factor 22		Hypothalami
	W.A.	Inniversal Alexant (MCIO), 55	growth factor	Midbrain
AK006679	1	h	1-	Conex
- WIND9/A	ł	Mus musculus adult male tesus cDNA, RIKEN full-length enriched library, clone 1700041N15 product (	eroush factor	Cortex
4 KW4	la		]	
4K005458	Prtpn	protectin-like protein N	growth factor	Striatum
				Striatum
AF210429	Pla2010	phosphotipase A2, group X	misceflaneous (secreted)	Midbrain

#### FIGURE 25A

#### DRUG TARGETING CANDIDATES

#### Colon Cancer

Accesion	Gen	Description	Class	
X15643	Adrb2	adrenergic receptor, beta 2		Region
NM_031257	Pickha2	pleckstrin homology domain-containing, family A (phosphoinosinde binding specific) member 2	signaling (receptor)	Cortex
NM_030733	Cipré3	G protein-coupled receptor PSP24-2 [Mus musculus] 100 %	signaling (receptor)	Hypothalamus
NM_030726	Cor90	G protein-coupled receptor 90	signaling (receptor)	Hypothalamus
		o process complete receptor 90	signaling (receptor)	Midbrain
1	ŀ		" " " " " " " " " " " " " " " " " " "	Cortex
	Ī		ì	
				Striatum
NM_030701	Puma-g	putative seven transmembrane spanning receptor purna-g		Hypothalamus
l			signaling (receptor)	Cortex
				Striatem
NM_030553	O1fr 160	Olfr160		Hypothalamus
NM_030258		hypothesical annual ACC 2014 A	signaling (receptor)	Hypothalamus
NM 022427	Gpr88	hypothetical protein, MGC:7035; hypothetical protein MGC7035 [Mus musculus] 100 %	signaling (receptor)	Hypothalamus
NM_021458	Fzd3	On browning complete secretary 89	signaling (receptor)	Hypothalamus
NM_021457	Fzd1	frizzled homolog 3 drosophils fzd3	signaling (receptor)	Hypothalamus
		frizzled-1	signaling (receptor)	Midbrain
NM_021406	Trem)	miggering receptor expressed on myeloid cells i		
NM_021368	Ors 16	odorant receptor 16	signaling (receptor)	Midbrain
	ł	i i	signaling (receptor)	Midbrain
	i		Ī	Cartes
NM_020598	01617	ol factory receptor 17		Hypothalamus
NM_020518	Ctal		signaling (receptor)	Midbrain
	F-'^'	cortical thymocyte receptor (X. laevis CTX) like	immune response (receptor)	Suiatum
h	۱			
NM_020515	Om 16	gene for odorant receptor A16	einaulina (assessa)	Hypothalamus
			signaling (receptor)	Cortex
NM_020488	Gabro	gaba-a receptor theta subunit family member	1	Striatum
I			signaling (receptor)	Midbrain
NM_020292	MOR32-4	olfactory receptor MOR32-4 - odorant receptor S46 gene (Mus musculus) 97 %		Cortex
_	1	7	miscellaneous (receptor)	Midbrain
NM_020288	Oral	administrative Ct		Hypothalamus
NM_019952	BsB	odorant receptor SI gene [Mus musculus] 100 %	miscellaneous (receptor)	Cores
NM_019691		B-cell stimulating factor	Signaling (receptor)	Cortex
	Gria4	glutamate receptor ionocropic ampa4 alpha 4 gris4	signaling (receptor)	
NM_019476	O16159	offictory receptor 159		Midbrain
i			signaling (receptor)	Midbrain
ł	1			Cortex
NM_018780	Strp5	secreted frizzled-related sequence protein 5		Hypothalamus
1 -	1	And a service seducate broken a	signaling (receptor)	Midbrain
1	Į.			Cortex
	!	1	1	Striatum
	1			
NM_016719	Grb14	growth factor receptor bound protein 16	signaling (receptor)	Hypothalamus
NM_016675	Cldn2	claudin 2		Cortex
l	ļ		signaling (receptor)	Cortex
NM_015766	Ebi	Epstein-Barr virus induced gene 3		Striatum
NM 013887	Opn4	opsin 4 (melanopsin)	signaling (receptor)	Hypothalamus
NM 013845	Rorl	opan - (neminpsin)	signaling (receptor)	Cortex
NM_013728	Olf: 154	receptor tyrosine kinase-like orphan ror I - ROR1_MOUSE Tyrosine-protein kinase transmembrane	signaling (receptor)	Striatum
		orisemy receptor 134	signaling (receptor)	Midbrain
NM_013619	Olfr67	olfactory receptor 67		
NM_013616	O16-64	offactory receptor 64	signaling (receptor)	Cortex
	1		signaling (receptor)	Cortex
NM_013483	Bontal	butyrophilin. subfamily I, member A1		Striatum
NM_011999	Cleasib	C-type (calcium dependent carbohydras area to a second	immune response (receptor)	Striatum
-	1	C-type (calcium dependent, carbohydrate recognition domain) lectin, superfamily member 6	signaling (receptor)	Midbrain
1	1			Cortex
l	l	1	1	Striatum
NM OUT	lea		1	Hypothalamus
NM_011519	Sde1	syndecan I	signaling (receptor)	
				Striatum
NM_011328	Sct	secretin	lat and the same of	Hypothalamus
NM_011216	Ptpro	protein tyrosine phosphatase, receptor type, O	signaling (receptor)	Cortex
	1		signating (receptor)	Cortex
NM_010483	Harsh	S. hadron promine (amount)	l	Striatum
NM_010327	Gp1bb	5-hydroxytryptamine (serotonin) receptor SB	signaling (receptor)	Midbrain
NM 009914	Cor3	glycoprotein lb, beta polypeptide	cell adhesion (receptor)	Cortex
NM_009651		chemokine (C-C) receptor )	immune response (receptor)	Midbrain
	Akap4	A kinase (PRKA) anchor protein 4	signaling (receptors	Cortex
NM_009493	V2r4	vomeronasal 2, receptor, 4	secution (secchion)	
NM_009325	Tbxa2r	thrombonane a2 receptor		Cartex
NM_008962	Ptgdr	prostaglandin D receptor	signating (receptor)	Striatum
NM_008720	Npcl	niemann pick type e1	Signaling (receptors	Contex
NM 008559	McIr	melanocoria I manus	signaling (receptor)	Midbrain
NA1_008479	Lag3	metanocortin I receptor	signaling (receptor)	Cortex
	-485	lymphocyte-activation gene 3	inumune response (receptor)	Midbrain
	1		and a second	Cortes
i	1		1	
1	l .		]	Striatum
NM_008364	lling	interleukin 1 receptor accessory protein		Hypothalamus
NM 008338	lfngr2	interferon gamma receptor 2	signaling (receptor)	dbram

#### FIGURE 25B

Accesion	Cen	Description	Class	
WM_008106	Opn I mw			Region
MM_00\$076	Gatur 2	RANGULA-RIMBOCALISMO and subsets recommend subsets about 3 1 1	signaling (receptor)	Cortex
NM_00\$074	Gabrg3	Extrata Subunit of the debasinha recessor	signating (receptor)	Cortex
250500 MY	Fzd4	frizzled homolog 4 drosophile frd4	signaling (receptor)	Hypothalamu
NM_007904	Ednth	endothelin-b receptor edurb	signaling (receptor)	Striatem
M_007900	Ect2		signaling (receptor)	Cortex
	1		signaling (receptor)	Midbrain
			3,,	Cornex
NM 007699	0			
	Chrm4	cholinergic receptur muscarinie 4 chrm4	signaling (receptor)	Striatum
NM_007680	Ephb6	and beauti	signaling (receptor)	Midbrain
M36699	Rho		signating (receptor)	Midbrein
AK017571	Chmb3		signaling (receptor)	Cortex
AK014671	i	antigen identified by monoclosed applicable bather over	Signaling (receptor)	Midbrain
	I	( I was a series ( I was a series)	signaling (receptor)	Midbrain
AK014543		progesterone membrane binding protein [73% Homo sapiens]		Cortex
4K010800	l .	Simpler to A40038 MHC close this common that	signaling (receptor)	Midbrain
	1.	Similar to A40038 MHC class I histocompatibility antigen H-2 M3 alpha chain procursor - mouse of	immune response (receptor)	Midbrain
	i			Striatum
AK009282	i .	RIKEN cDNA 1310010M24 gene		Hypothalamus
AK006680		CLD2_MOUSE CLAUDIN-2 23 %	signaling (receptor)	Hypothalamus
	1		signaling (receptor)	Correx
AF320126	Gmil			Striansm
AF282301	MOR224-4	metabotropic glutamate receptor type 1 (Glur1)	Signating (receptor)	Hypothalamu
AF132300	mor17-1		signaling (receptor)	Corres
A 122300	mari /-:		signaing (receptor)	Conex
			Suring (.cochice)	Striatum

#### Breast Cancer

(59150	Gen	Description	Class	Region
	Tcrb-V20	T-cell receptor beta, variable V20	signaling (receptor)	Midbrain
189915	l		greating (receptor)	
	Filt	F11 receptor	simulian (manage)	Hypothalamu
82439	Ptprn2	protein tyrosine phosphatase 13-2beta ptp an autoantigen in insulin-dependent diabetes mellinus; ph	signaling (receptor)	Midhrain
(M_033269	Chrm3	ACM3_MOUSE Muscarinic acetylcholine receptor M3 (Mm3 mAChR) 100 %/		Cortex
			signaling (receptor)	Cortex
NM_032465	Tactile	T cell activation, increased late expression		Hypothalamu
NM_032400	Carsi	G protein-coupled receptor 91	signaling (receptor)	Hypothalamu
VM_032002	Nrg4	neuregulin 4 nrg4	signaling (receptor)	Hypothalamu
			Signaling (receptor)	Midbrain
				Cortex
NM_031373	Ogt	opicid growth factor receptor		Hypothalamu
	-		signaling (receptor)	Michrain
NM 030735	V3R9	pheromone receptor V3R9		Hypoth alamu
-		productive vigo	signaling (receptor)	Midbrain
	1			Cortex
NM_030701	Puma-g	Duraning and a second of the s		Hypothalams
	· wita-g	putative seven transmembrane spanning receptor puma-g	signaling (receptor)	Midbrain
NM_030553	O16 160	016160		Hypothalamu
NM_028595	Ms4a6c		signaling (receptor)	Hypothalamu
(11,02037)	uradaoc .	membrane-spanning 4-domains, subfamily A, member 6C	immune response (receptor)	Cortex
NM_025771	Cumap2	contactin associated protein-like 2	cell adhesion (receptor)	Hypothalamu
	1.		cen suresion (receptor)	Cortex
NM_021458	Fzd3	frizzled homolog 3 drosophila fzd3	-1	Hypothalama
NM_021457	Fzdi	frizzled-1	signaling (receptor)	Hypothalamu
NM_021406	Treml	triggering receptor expressed on myeloid cells 1	signaling (receptor)	Cortex
NM_021332	Glp 1 r	glucagon-like peptide I receptor	signaling (receptor)	Cortex
NM_021325	Mox2r	antigen identified by monoclonal antibody MRC OX-2 recepto	signaling (receptor)	Midbrain
NM_020598	Olfr 17	olfactory receptor 17	signaling (receptor)	Hypothalamu
NM_020515	Ora 16	gene for odorani receptor A16	signating (receptor)	Midbrain
		James de la constitut Alo	signaling (receptor)	Midbrain
NM 020291	Ors25	odorant receptor \$25		Hypothalamu
NM 020290	075.18	odorant receptor \$18 gene	signaling (receptor)	Midbrain
NM_020288	Orsi		Signaling (receptor)	Hypothalamu
NM_020278	Leil	odorant receptor S1 gene (Mus museulus) 100 %	miscellaneous (receptor)	Hypothalamu
NM_019691	Gris4	leucine-rich repeat LGI family, member !	signating (recentor)	Hypothalarma
NM_019583	(1) 7br	glutamate receptor ionograpic ampa4 alpha 4 gria4	signating (receptor)	Hypothalamu
		interleukin 17B receptor	signaling (receptor)	Cortex
M_019485	Q16-70	olfactory receptor 70	signaling (receptor)	
	l l		arguming (receptor)	Cortex
NM_019421	1.	hypothetical protein 425Q18-1	-ershelia	Hypothalamu
NM_018766	Ntsr	neurotensia receptor	metabolism (receptor)	Hypothalamu
NM_017480	icos	inducible T-cell co-stimulator	signaling (receptor)	Hypothalamo
		i	immune response (receptor)	Midbrain
NM_016719	Grb14	growth factor receptor bound protein 14		Hypothalame
NM_016696	Gpc I	glypican I	signaling (receptor)	Midbrein
NM_015743	Nr4a3	nuclear receptor subfamily 4, group A, member 3	signaling (receptor)	Midbrain
NM_013845	Rorl	Interpret Comment to the property of the property	transcription (receptor)	Midbrain
SM 013728	Olf:154	receptor tyrosine kinase-like orphan ror I - ROR1_MOUSE Tyrosine-protein kinase transmembrane offactory receptor 154	signaling (receptor)	Hypothalarm
-		,,,	signaling (receptor)	Midbrain
VM_013662	Semabb	sema domain programme and a transport	a Water	Hypothalama
M 013658	Senuta	sema domaia, transmembrane domain (TM), and eytoplasmic domain, (somaphorin) 6B	signating (receptor)	Hypothalarm
M 013622	Oprdi	sems domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short exercises and	signating (receptor)	Cortex
VM_013619	Olfre7		signaling (receptor)	Hypothalamu
	Olifo	olfactory receptor 67	signaling (receptor)	Midbrain
	1			
				Cortex
NM_013618	O1666	olfactory receptor 66	-Itit	Hypothalamu
NM 012584	Life	leukemia inhibitory factor receptor	signaling (receptor)	Hypothalamu
	-		signaling (receptor)	Hypothalamu

#### FIGURE 25C

SM_013476	Buntal	buryrophilin, subfamily I, member Al	Chn	Region
		The second of th	immune response (receptor)	Midbrain
34 011174				Cortex
	l <sub>A</sub> r	androgen recept		Hypothalama
M_011976	Sema4g	tems downer amountable to the	transcription (receptor)	Midbrain
M 011798	Xort	serua domain, immunoglobulin domain (lg), transmembrane domain (TM) and short cytoplasmic of chemokine (C motif) receptor 1	(signaling (receptor)	Hypothalami
	P	ementioning (c. mineri) secuptor (	signating (receptor)	Midbrain
M_011519	Sdel	nature 4		Hypothalam
M_011216	Ptpro	syndecan 1	signaling (receptor)	
M_010482		protein syrosine photobause, receptor type, O	signaling (receptor)	Hypothalam
	Hurth	5-hydroxyeryptamine (serotonin) receptor 1B	signaling (receptor)	Hypothalam
	1		signaming (receptor)	Midbrain
	1		1	Cortex
M_010346	Grb7	growth factor receptor bound protein 7 clone mgc:5653; epidermal receptor-binding grb7	l	Hypothalam
M_010338	Cpr37	TO Protesti-Company recentor 37	growth factor (receptor)	Hypothalam:
M_010317	Gng4	guanine nucleotide binding protein (G protein), gamma 4 subunit	signaling (receptor)	Hypothalam
M_D10254	Catr2	galania receptor 2	signaling (receptor)	Michrain
			signaling (receptor)	Midbrain
M_010077	Drd2	doparnine receptor 2		Hypothalam
M_009912	Cat	chemokine (C-C motif) receptor 1	signaling (receptor)	Hypothalam
M_009903	Cldn4	claudin 4	Signating (receptor)	Midbrain
M_009886	Celui		signaling (receptor)	Hypothalam
M_009827	Cokar	cadherin egf lag seven-pass g-type receptor cels: I cholecystokinin A receptor	cell adhesion (receptor)	Midbrain
		Connect American V Lecebras	signaling (receptor)	Midbrain
M_009602	Chrob2	Address to the second	1	Hypothatam
007002	Curanz	cholinergic receptor, nicotinic, beta polypeptide 2 (neuronal)	signaling (receptor)	nypomaram
M 009521		i e	Tracepus,	Midbrain
	Wni3	wingless-related MMTV integration site 3	Signation (comme	Hypothalam
M_009493	V2r4	vomeronasal 2, receptor, 4	signaling (receptor)	Cortex
	L .		receptor	Midbrain
M_009370	Tgfbri	transforming growth factor beta receptor )	l	Hypothalam
M_00929D	Wntla	wingless-related MMTV integration site 8A	signaling (receptor)	Hypothalam
	Rrh	retinal pigment epithelium derived rhodopsin homolog	signaling (receptor)	Hypothalam
	Ptgdr	prostaglandin D receptor	signaling (receptor)	Cortex
	Preh2	patched homolog 2	signaling (receptor)	Midbrain
M 008938	Rds		(signaling (receptor)	Cortex
-		retinal degeneration, slow (retinitis pigmentosa 7)	cell adhesion (receptor)	Cortex
M_008746	Nork	parametric a such at	1	Hypothalama
M_008720	Npcl	neurotrophic tyrosine kinase, receptor, type 3	signaling (receptor)	Hypothalama
M 008716	Notch3	nicmann pick type cl	signaling (receptor)	Midbrain
		Notch gene homolog 3 (Drosophila)	signaling (receptor)	
•	Nmbr	neuromedia B receptor	signaling (receptor)	Hypothalams
	Mc3r	melanocortin recuptor		Cortex
M_008533	Ly78	lymphocyte antigen 78 ly78	signaling (receptor)	Hypothatamu
MI_008479	Lag3	lymphocyte-activation gene 3	immune response (receptor)	Hypothalami
M_008353	1112761	interleukin 12 receptor, beta !	immune response (receptor)	Hypothalams
M_008348	III Ora	interleukin-10 receptor alpha	signaling (receptor)	Cortex
		, , , , , ,	signaling (receptor)	Midbrain
				Cortex
M_008338	lfngr2	interferon gamma receptor 2		Hypothalamy
M_008315	Her?	5-hydroxycryptamine (serotonin) receptor 7	immune response (receptor)	Midbrain
	Hirla	Charles and properties (receptor)	signaling (receptor)	Hypothalams
	Opsimw	S-hydroxytryptamine serotonin receptor 1a htr1a	signaling (receptor)	Hypothalame
	Gabri 2	opsin I (cone pigments), medium-wave-sensitive (color blindness, deutan)	signaling (receptor)	Cortex
M 008074	Gabry 3	gamena-arrangoutyric acid gaba-a receptor subunit the 2 exhre?	signaling (receptor)	
M 008070		gamma subunit of the gaba-alpha receptor	signaling (receptor)	Hypothalamo
	Gabrb2	gamuna-aminobutyric acid gaba-a receptor subunit beta 2 gabrb2		Hypothalamu
M_008069	Gabrb 1	Seriona-arminocutyric acid (GABA-A) recenter, subspirit here #	signaling (receptor)	Hypothalamu
M_008055	Fzd4	jurazzed nomolog 4 drosophila fzd4	signaling (receptor)	Hypothalamu
	Dixi	[deltex]	signaling (receptor)	Midbrain
	Ephz8	Eph receptor A8	signaling (receptor)	Cortex
	Ednito	endothelin-b receptor edorb	signaling (receptor)	Hypothalama
	Ea2	ect2 ancagene	signaling (receptor)	Hypothalamu
	Don	decorin	signaling (receptor)	Hypothelam
M_007407	Adcyaptri	adenylate cyclase activating polypeptide 1 receptor 1	Signaling (receptor)	Cortex
(34)7)	Спу	complement receptor related protein	Signaling (receptor)	Midbrain
26157	Pigir	prostacyclin receptor	immune response (receptor)	Cortex
	1	L	signaling (receptor)	Midbrein
C004794	Fragi	FGF promine agriculture accepts to		Cortex
K019508		FGF receptor activating protein 1	Signating (receptor)	Hypothalama
K017036		cadherin EGF LAG seven-pass G-type receptor 1, cadherin EGF LAG seven-pass G-type receptor (	signaling (receptor)	Hypothalana
K014543			signaling (receptor)	Hypothalama
	Sreb3	progesterone membrane binding protein [73% Homo sapiens]	signaling (receptor)	
		SKERS	signaling (receptor)	Hypothalama
NO12604	P2ry12	purinergic receptor P2Y. G-protein coupled 12	signaling (receptor)	Hypothalam
	1	Similar to type I transmembrane receptor (seizure-related protein) (17% Home series)	signaling (receptor)	Midbrain
	l	IMPERIORIA SOLUCI IMUS BENE		Hypothalana
K012717	1	Similar to A40038 MWC along I have not the	signating (receptor)	Cortex
K012717	I	- mages are repaired arising procursor - mouse	immune response (receptor)	Cortex
K012717 K010800				Hypothalann
K012717 K010800	l	157963 natriuretic pentide recentor A - monte 50 3 4 7		
K013276 K012717 K010800 K010727 K009282		157963 natriuretic peptide receptor A - mouse 50,36 %	Signaling (receptor)	Midbrain
K012717 K010800 K010727 K009282	Cloud	RIKEN cDNA 2310010M24 gene	Signaling (receptor) signaling (receptor)	Midbrain
K012717 K010800 K010727 K009282	Class	157963 natriuretic peptide receptor A - mouse 50,36 % RIKEN cDNA 2310010M24 gene chloride channel 1	signaling (receptor)	Midbrain
K012717 K010800 K010727 K009282 J011106		RIBEN CONA 2310010M24 genz chloride channel I		Midbrain Midbrain
K012717 K010800 K010727 K009282 J011106	Clan I MOR224-6	RUSEN CDNA 2310010M24 gene chloride channel I colfactory receptor MOR224-6	signaling (receptor) transport/receptor	Midbrain Midbrain Hypothalami
K012717 K010800 K010727 K009282 J011106 F282302 F282300	MOR224-6	RUEEN CDNA 2310010M24 gene chloride channel 1  clofactory receptor MOR224-6  Mus musculus odorant recentor M15 sene	signaling (receptor) transport/receptor signaling (receptor)	Midbrain Midbrain Hypothalams Hypothalams
K012717 K010800 K010727 K009282 J011106 F282302 F282300 F282291	MOR224-6	RUEEN CDNA 2310010M24 gene chloride channel 1  clofactory receptor MOR224-6  Mus musculus odorant recentor M15 sene	signaling (receptor) transport/receptor signaling (receptor) signaling (receptor)	Midbrain Midbrain Hypothalami Hypothalami Midbrain
K012717 K010800 K010727 K009282 J011106 F282302 F282300 F282291 F123300	MOR224-6	RUEEN CDNA 2310010M24 gene chloride channel I  Olfactory receptor MOR224-6  Mus musculus odorant receptor M15 gene olfactory receptor MOR171-8	signaling (receptor) transport/receptor signaling (receptor) signaling (receptor) signaling (receptor)	Midbrain Midbrain Hypothalamu Hypothalamu
K012717 K010800 K010727 K009282 J011106 F282302 F282300 F282291 F123300	MOR224-6	KIREN CDNA 2310010M24 gene chloride channel 1  offactory receptor MOR224-6  Mus musculus odorant receptor M15 gene offactory receptor MOR171-8  offactory receptor MOR171-8	signaling (receptor) transport/receptor  signaling (receptor) signaling (receptor) signaling (receptor) signaling (receptor)	Midbrain Midbrain Hypothalamu Hypothalamu Midbrain
K012717 K010800 K010727 K009282 J011106 F282302 F282300 F282291 F123300	MOR224-6 MOR173-8 mor17-1	RUEEN CDNA 2310010M24 gene chloride channel I  Olfactory receptor MOR224-6  Mus musculus odorant receptor M15 gene olfactory receptor MOR171-8	signaling (receptor) transport/receptor signaling (receptor) signaling (receptor) signaling (receptor)	Midbrain Midbrain Hypothalamu Hypothalamu Midbrain Cortex

## FIGURE 25D

## Lung Cancer

X51552 UR2439 NM_053094 NM_033269	Terb-V20	Description  [F-cell receptor beta, variable V20	Class	Region
UR2439 NM_053094		·	signaling (receptor)	Corres
NM_053094	Fgf6	fibroblast growth freme 6		Striatum
	Ptpm2	protein tyrosine phosphatase is-libets pip an autoantigen in insulin-dependent diabetes mellitus, ph	signaling (receptor)	Cones
	C4163		(Signaling (receptor)	Midbrain
	Chmi	CD163 antigen	signaling (receptor)	Cortes Midbrain
		ACM3_MOUSE Musearmic acetylcholine receptor M3 (Mm3 mAChR) (00 % /	signaling (receptor)	Midbrain
NM_032400	Gpr91	G protein-coupled receptor 91		Cortes
			signaling (receptor)	Midbrain
NM_032002	Nrg4	acureguim 4 arg4	Summahana dananananan	Corses
W_031879	Phop	laterate to the state of the st	Signating (receptor)	Midbram Corex
NM_031373	Ozfi	pleckstrin homology domain interacting protein opioid growth factor receptor	signaling (receptor)	Midbrain
-		- Providence of the second	signaling (receptor)	Midbram
	1			Contra
			l	Stratum
NM_031257	Pickha?	pleckstrin homology domain-containing, family A (phosphotocitide binding specific) member 2	diameters (community	Hypothalamus
NM_030735 NM_030726	V3R0 Gpr90	President stocking 43K3	signaling (receptor) signaling (receptor)	Cortex
M_030781	Puma-g	G protein-coupled receptor 90	signaling (receptor)	Hypothalamus Striatum
	· u.u.a-g	putative seven transmembrane spanning receptor pama-g	signahng (receptor)	Cores
				Striatum
NM_030553	O18 160	Olfr160	į	Hypothalamus
NM_030258		hypothetical protein, MGC:7035; hypothetical protein MGC7035 [Mus musculus] 100 %	signaling (receptor)	Stratum
M_029926	Irak4	Intersection   1675-beth-120013100 Kinate 4	signaling (receptor)	Midbrain
NM_028595	Ms4a6c	membrane-spanning 4-domains, subfamily A, member 6C	immune response (receptor)	Hypothalamus
NM_025771	Commen		mmune response (receptor)	Cortex
	Cntnap2	contactin associated protein-like 2	cell adhesion (receptor)	Hypothalamus Midbrain
NM 022427	Gpr88	Garwen muded		Cortex
NM_021458	Fzd3	G-protein coupled receptor 88 friezled homolog 3 drosophila fad3	signaling (receptor)	Midbrain
M_021407	Trem3	inggening receptor expressed on myeloid cells 3	signaling (receptor)	Hypothalamus
-		mesoning reches criticated on mixtond Colle 3	signaling (receptor)	Sinatum
AW_051398	Ors 16	odorani receptor 16		Midbrain
NM_021364	Cleast5	C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 5	signaling (receptor)	Hypothalamus
VM_021332	Glplr	Processor, and behalfe I (CEASTER)	immune response receptor?	Midbrain
NM_021325	Mon2r	antigen identified by monoclonal antibody MRC OX-2 recepto	signaling (receptor)	Striatum
			signaling (receptor)	Midbrain
NM_021319	Palyrpi		1	Corex
NM_020515	Orale	peptidoglycan recognition protein-like	signaling (receptor)	Hypothalamus Hypothalamus
VM 020292	MORJ2-4	gene for odorani receptor A16	signaling (receptor)	Hypothalamus
VM_019923	ltpr2	olfactory receptor MOR32-4 - odorant receptor \$46 gene [Mus musculus] 97 % anositol 1.4.5-triphosphate receptor 2	misoclianeous (receptor)	Cortes
_		and and and an inches and a second a	signaling (receptor)	Midbrain
M_019691	Gris4	ghtamate receptor ionotropic ampa4 alpha 4 gris4		Hypothalamus
4M_019583	1117br	interleaken 178 receptor	signaling (receptor)	Cortex
/14 m10.00			signaling (receptor)	Midbrain
VM_019476	Otf: 159	olfactory receptor 159	signating (receptor)	Cortex
			and the state of t	Midbrain Conex
M_018780	Sfrp5	secreted frizzled-related sequence protein 5		Hypothalamus
M_016873	Wisp2	WNT1 inducible signaling pathway protein 2	signaling (receptor)	Conex
969910 Wi	Gpc i	ghpican I	signaling (receptor)	Hypothalamus
			signaling (receptor)	Cortex
M_015743	Nr4a3	muclear receptor subfamily 4, group A, member 3		Stratura
NM_015738	Galr3	galanin recuptor 3	transcription (receptor) signaling (receptor)	Cortex
		$\Lambda$	signating (receptor)	Midbrain Striatum
VM 015732	Axin2	axin2		Hypothalamos
VM_013845	Rori		Signaling (neceptor)	Cortex
_		receptor tyrosine kinase-like orphan ror I - RORI_MOUSE Tyrosine-protein kinase transmembrane	ugnaling (receptor)	Midbrain
M_013730	Slam	signating lymphocyte activation molecule		Hypothalamus
M_013728	O1fr154	olfactory receptor 154	signaling (receptor)	Hypothalamus
	_		signaling (receptor)	Midbrain
M_013662	Sema6b	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6B	signation (a.e)	Cortex
NM 013658	Sema4a Olfr67	seria surrain, immunoglobulin domain (lg), transmembrane domain (TM) and shore excellentia di	signaling (receptor)	Striatum Striatum
	J. 1007	olfactory receptor 67	signaling (receptor)	Cortex
	Olfren			Stratum
4M_013619			signaling (receptor)	
4M_013619 4M_013618	Life	olfactory receptor 66	manning (reschoot)	IM/POUISTABlus
iM_013619 iM_013618 iM_013584	Life Fprl	leukemia inhibitory factor receptor	signaling (receptor)	Conus
IM_013619 IM_013618 IM_013584 IM_013521		leukema inhibitory factor receptor formyt peptide receptor I	signaling (receptor) signaling (receptor)	Cones Stratum
IM_013619 IM_013618 IM_013584 IM_013521	Fpr I	leukemia inhibitory factor receptor	signaling (receptor)	Cortes Stratum Midbrain
IM_013619 IM_013618 IM_013584 IM_013521	Fpr I	leukema inhibitory factor receptor formyt peptide receptor I	signaling (receptor) signaling (receptor)	Concy Stratum Midbrain Concy
iM_013619 iM_013618 iM_013584 iM_013521 iM_013483	Fpri Binta)	leukema inhibitory factor reseptor formyl peptide receptor I butyrophiha, subfamily I, member A1	signaling (receptor) signaling (receptor)	Cortex Striatum Midbrain Cortex Striatum
VM_013619 VM_013618 VM_013584 VM_013521	Fpr I	leukemia inhibitory factor receptor formyl peptide receptor I bulyrophilin, subfamily i, member A1 androgen recept	signaling (receptor) signaling (receptor) immune response (receptor)	Cortex Striatum Midbrain Cortex Striatum Hypothalamus
VM_013618 VM_013584 VM_013521 VM_013483	Fpri Binta) Ar	leukemia inhibitory factor roseptor formyl peptide receptor I buly rophilin, subfamily I, member AI androgen rosept	signaling (receptor) signaling (receptor)	Striatum Midbrain Cortex Striatum Hypothalamus Cortex
IM_013619 IM_013618 IM_013584 IM_013521 IM_013483 IM_013476 IM_013476	Fpri Binta} Ar W:fi	leukemia inhibitory factor receptor formyl peptide receptor I butyrophilina subfamily I member A1 androgen recept Wat inhibitory factor I	agnaling (receptor) signaling (receptor) immune response (receptor) transcription (receptor)	Contex Striatum Midbrain Contex Striatum Hypothalamus Contex Striatum
M_013619 M_013618 M_013584 M_013521 M_013483	Fpri Binta) Ar	leukemia inhibitory factor roseptor formyl peptide receptor I buly rophilin, subfamily I, member AI androgen rosept	agnaling (receptor) agnaling (receptor) immuse response (receptor)  transcruption (receptor) asgnaling (receptor)	Cortex Striatum Midbraia Cortex Striatum Hypothalamus Cortex Striatum Striatum
VM_013619 VM_013618 VM_013584 VM_013521 VM_013483 VM_013476 VM_013476	Fpri Binta} Ar W:fi	leukemia inhibitory factor receptor formyl peptide receptor I butyrophilina subfamily I member A1 androgen recept Wat inhibitory factor I	agnaling (receptor) signaling (receptor) immune response (receptor) transcription (receptor)	Cortex Striatum Midbrata Cortex Striatum Hypothalamus Cortex Striatum Striatum Midbrata
IM_013619 IM_013618 IM_013584 IM_013521 IM_013483 IM_013476 IM_013476 IM_011415 IM_011798	Fpri Binta} Ar W:fi	leukema inhibitory factor receptor formyl pepade receptor I butyrophiha, subfamily I, member AI  androgen recept  Wat inhibitory, factor I chemok me (C motif) receptor 1	agnaling (receptor) agnaling (receptor) immuse response (receptor)  transcruption (receptor) asgnaling (receptor)	Cortex Striatum Midbraia Cortex Striatum Hypothalamus Cortex Striatum Striatum
IM_013619  IM_013618  IM_013584  IM_013521  IM_013483  IM_013476  IM_013476  IM_011915  IM_011798	Fpri Binta) Ar Wifi Xeri	leukemia inhibitory factor receptor formyl peptide receptor I butyrophilina subfamily I member A1 androgen recept Wat inhibitory factor I	agnaling (receptor) agnaling (receptor) immuse response (receptor)  transcruption (receptor) asgnaling (receptor)	Cortex Stratum Midbraia Cortex Stratum Hypothalamus Cortex Stratum Stratum Stratum Midbraia Cortex
IM_013619  IM_013618  IM_013584  IM_013521  IM_013483  IM_013476  IM_013476  IM_011798  IM_011776  IM_011776  IM_011776	Fpri Binta) Ar Wifi Xeri	leukema inhibitory factor receptor formyl peptide receptor I butyrophilin, subfamily I, member A1  androgen recept Wat inhibitory factor I chemokine (C mouf) receptor I  zona pellucida glycoprotein 3	agnaling (receptor) signaling (receptor) immune response (receptor) transcription (receptor) signaling (receptor) signaling (receptor)	Cortex Striatum Midbrain Cortex Striatum Hypothalamus Cortex Striatum Striatum Midbrain Cortex Striatum Striatum Striatum
M_013619  SM_013514  SM_013514  SM_013517  SM_013413  SM_013476  SM_011742  SM_011776  SM_011776  SM_011812	Fpri Binla) Ar Wifi Xeri Zp3 Hirlb Gnal2	leukemia inhibitory factor receptor formyl peptide receptor 1 buty rophilin, subfamily 1, member A1  androgen recept  Wat inhibitory factor 1 chemokime (C moul) receptor 1  zona pellucida glycoprotein 3  5-bydroxytyptamine (sercotomia) receptor [B] guanten nucleotide bindhar protein, sloba ) ?	ingnaling (receptor) signaling (receptor) immuse extronal (receptor)  transcription (receptor) signaling (receptor) signaling (receptor) signaling (receptor) signaling (receptor)	Cortex Striatum Middraia Cortex Striatum Hypothalamus Cortex Striatum Striatum Middrain Cortex Striatum Striatum Striatum Striatum Striatum Striatum Striatum Striatum Striatum
IM_013619 IM_013618 IM_013584 IM_013581 IM_013483 IM_013483 IM_013476 IM_011915 IM_011778	Fpri Binla) Ar Wifi Xeri Zp3 Htrib Goal 2 Chadr	leukemia inhibitory factor receptor formyl peptide receptor 1 buty rophilin, subfamily 1, member A1  androgen recept  Wat inhibitory factor 1 chemokime (C moul) receptor 1  zona pellucida glycoprotein 3  5-bydroxytyptamine (sercotomia) receptor [B] guanten nucleotide bindhar protein, sloba ) ?	agnaling (receptor) agnaling (receptor) immuse response (receptor)  transcription (receptor) agnaling (receptor) agnaling (receptor) agnaling (receptor) agnaling (receptor)	Cortex Striatum Midbraia Cortex Striatum Hypothalamus Cortex Striatum Striatum Midbraia Cortex Striatum Striatum Striatum Striatum Striatum Hypothalamus Midbraia
M_013619  M_013618  M_013544  M_013544  M_013548  M_01348  M_01348  M_013476  M_013476  M_011915  M_011798  M_011798  M_011798  M_011798  M_011798  M_011798  M_011798	Fpri Binlai Ar Wifi Xeri Zp3 Htrib Gnai 2 Chadr Ccr3	leukemia inhibitory factor receptor formyl peptide receptor 1 buly rophilin, subfamily 1, member A1  androgen recept  Wat inhibitory factor 1 chemokime (C motif) receptor 1  zona pellucida glycoprotein 3  S-bydroxytryptamine (acrotomia) receptor (B guantine nucleotide binding protein, alpha 12 covaschessivus and ademosirus neceptor chemokime (C-C) receptor 3	agnaling (receptor) agnaling (receptor) immune response (receptor)  transcruption (receptor) agnaling (receptor) agnaling (receptor)  agnaling (receptor)  agnaling (receptor) agnaling (receptor) agnaling (receptor)	Cortex Striatum Middrain Cortex Striatum Hypothalamus Cortex Striatum Striatum Striatum Striatum Striatum Hypothalamus Striatum Hypothalamus Middrain Hypothalamus Striatum Striatum
VM_013619 VM_013618 VM_013584 VM_013521 VM_013483 VM_013476 VM_013476	Fpri Binla) Ar Wifi Xeri Zp3 Htrib Goal 2 Chadr	leukemia inhibitory factor receptor formyl pepnde receptor I buryrophilin, subfamily I, member A1  androgen recept  Wat inhibitory factor I chemokine (C mouf) receptor I  aona pellucida glycoprotein 3  5-bydroxytyptamine (serotomia) receptor IB guaniae nucleotide binding protein alpha 12 covacekeur san adadonostrus neceptor	agnaling (receptor) agnaling (receptor) immuse response (receptor)  transcription (receptor) agnaling (receptor) agnaling (receptor) agnaling (receptor) agnaling (receptor)	Cortex Striatum Midbraia Cortex Striatum Hypothalamus Cortex Striatum Striatum Midbraia Cortex Striatum Striatum Striatum Striatum Striatum Hypothalamus Midbraia

#### FIGURE 25E

Accesion	Gen	Description 1 TOOR 251		
NM 009854	Cd7	cd7 saniero	Ctass	Region
NM 009827	Cdu	cholecystokinin A receptor	immune response (receptor)	Hypothalamus
i			Signaling (receptor)	Cortex
NM_009524	WatSa	wingless-related MMTV integration site SA		Striatum
NM_009504	VΦ	vitamia d receptor	signaling (receptor)	Hypothalamus
NM_009493	V2r4	vomeronasal 2, receptor, 4	signaling (receptor)	Midbrain
NM_008962	Prgdr	prostaglandin D receptor	receptor	Hypothalamus
NM_008938	Rds	retinal degeneration, slow (retinitis pigmentosa 7)	signaling (receptor)	Cortex
NM_008746	Nerk3	neurocophie syrosine kinese, receptor, type 3	cell adhesion (receptor)	Hypothalarms
-			signaling (receptor)	Cortex
NM_008716	Notch3	Notch gene homolog 3 (Drosophila)		Hypothalamus
	1	Pare nomined a (Diesobuita)	signating (receptor)	Midbrain
	1			Striatum
NM_008703	North	newomedin B receptor	h	Hypothalamus
NM 008559	Melr	metanecortin I receptor	signating (receptor)	Hypothalamus
NM 008512	Lpl	low density lipoprotein receptor-related protein t	signating (receptor)	Hypothalamus
NM_008479	Lag3	lymphocyte-activation gene 3	signaling (receptor)	Midbrain
-	-	-77	inumune response (receptor)	Midbrain
NM_003401	Itgan	integrin alpha M	, , , , , , , , , , , , , , , , , , , ,	Hypothalamus
NM 001398	Itga7	integrin slpha 7	cell adhesion (receptor)	Hypothalamus
NM_008364	litrap		cell adhesion (receptor)	Stristum
		interleukin   receptor accessory protein	signaling (receptor)	Cortex
NM_008353	(1) 2rb1	insuladia 19	, , , , , ,	Hypothalamus
•		interleukin 12 receptor, beta 1	signaling (receptor)	Midbrain
	[			Cortex
NM_008348	III Ora	investments 10 annual to		Hypothalamus
		interleukin-10 receptor alpha	signaling (receptor)	Cortex
	1			Striatum
NM_008338	lfngr2	invertina and a second		Hypothalamus
NM_008285	Hrb1	interferon gammu receptor 2	insmune response (receptor)	Hypothalamus
_000205	13741	histarnine receptor H 1	signaling (receptor)	Cortex
NM_008172	Grin2d		( and )	Hypothalarms
NM_008166	Gridt	glutamaze receptor channel subunit epsilon 4	signaling (receptor)	Striatum
NM 008106		glutarrate receptor ionograpic delta grid l	signaling (receptor)	
NM_008070	Opn tmw Gabrb2	opsin I (cone pigments), medium-wave-sensitive (color blindness, deutan)	signaling (receptor)	Hypothalamus Hypothalamus
NM_008069		[Earturns-Ettilin COUNTIC acid gaba-a recentor turbunit here 2 cobob?	signaling (receptor)	
NM_008051	Gabrol	[Bitturia-arminobutyric acid (GABA-A) recentor, subunit bers 1	signaling (receptor)	Cortex
NM 008055	Fzd8 Fzd4	parative transmending receptor frizzled \$	signaling (receptor)	Hypothelamus Midbrain
NM 007975	F2rt3	frizzled homolog 4 drosophila fzd4	signaling (receptor)	Cortex
NN_007975		protease-activated receptor 4 par4 g protein-coupled receptor thrombin	signaling (receptor)	Midbrain
NM_007939	F2rt3	protesse-activated receptor 4 par4 g protein-coupled receptor thrombin	signaling (receptor)	Hypothalamus
NM 007904	Epha8	Epit receptor As	signaling (receptor)	
NM 007779	Edarb	endothetin-b receptor edurb		Cortex
NM_007699	Csflr Chrm4	e-fins proto-oncogene protein precursor as -19 to 957; adult male liver riken cons cione: 130000xn2	Signating (receptar)	Hypothalarrus
			signaling (receptor)	Hypothalamus Midbrain
NM_007412 M36699	Admy	adrenomedullin receptor	signaling (receptor)	
L24495	Rho	rhodopsia	signaling (receptor)	Cortex
L07379	Tnfrsf7	turnor necrosis factor receptor superfamily, member 7	Signating (receptor)	Cortex
	Ghrhr	(clone porn?) growth hormone-releasing factor recentor	rignaling (receptor)	Striatum
BC016104 BC007481	Gpr108	C protein-coupled receptor 108		Hypothalamus
	Planb2	plexin B2	Signaling (receptor)	Striatum
AK017036			Signaling (receptor)	Striatum
AK016763		RIKEN cDNA 4933411803 gene	signaling (receptor)	Striatum
AK015192	1	interleukin 6 receptor, alpha: 11-6 receptor sinha chair 16.79/ Mars and the	signaling (receptor) ?	Striatum
AK014671	l .	antigen identified by monoclonal antibody MRC OX-2 receptor (41% Mus musculus)	Signaling (receptor)	Striatum
	1	•	signaling (receptor)	Midbrain
AK014543	i	progesterone membrane binding protein [73% Homo sapiens]		Hypothalamus
	1		signaling (receptor)	Midbrain
	L			Cortex
AK014326	Sreb3	SREB3	l	Hypothalarnus
AK013804	P2ry12	purinergic receptor P2Y, G-protein coupled 12	signaling (receptor)	Hypothalamus
AK011967	P2y5	[purimergic receptor (family A group 5)	signaling (receptor)	Striatum
AK010800	1	Similar to A40031 MMC class 1 histonymusikiling	signaling (receptor)	Hypothalamus
			immune response (receptor)	Midbrain
AK010720	1	Similar to seven transmembrane protein TM7SF3 [82% Homo sapiens]		Hypothalamus
AK009282		RIKEN cDNA 2310010M24 gene	signaling (receptor)	Hypothalamus
	ŀ		signaling (receptor)	Midbrain
	1			Cortex
AK004472	1	polycychemia rubra vera 1; cell surface receptor; cell surface receptor [Homo sapiens] 49 %	l	Hypothalamus
	1	49 %	signaling (receptor)	Midbrain
AJ011106	Clent	chloride channel )		Hypothalamus
AF282302	MOR224-6	olfactory receptor MOR224-6	transport/receptor	Midbrain
AF282301	MOR224-4	olfactory receptor MOR224-4	signaling (receptor)	Midbrain
	1	The state of the s	signaling (receptor)	Midbrain
AF282291	MOR171-8	olfactory receptor MOR171-8	l	Hypothalannus
AF282286		Mus musculus odorant receptor K30 gene	signaling (receptor)	Cortex
AF133300	mor (7-)	olfatory receptor mor17-1	signating (receptor)	Cortex
		Amen's control (1921-1-1)	signaing (receptor)	Cortex
AF056187	tgftr	intulinable grouph factor i accounts in 6:		Striatum
- /	J.****	insulin-like growth factor i receptor igf-i	signating (receptor)	Striatum
AF045766	Ger33	comban o puestio assult de constante de la con		Hypothalamus
AF027131	Muc3	orphan g protein-coupled receptor gpr33 related to chemoattractant receptors mucia 3, intestinal	signaling (receptor)	Hypothalanius
	,	Linde in N. mor Hills.	signaling (receptor)	Cones

## FIGURE 26

# CNS markers differentially expressed in all tumors analyzed

Accession	Gen	Description	le		
X72307	Hgf	hepatocyte growth factor	Class	Temor	Region
		in-hanning Broads Mctor	growth factor	Colon	Cortex
			1	Breast	Midbrain
	1				Corner
	ļ		l.	Lung	Midhrain
	1		i	1 -	Cortex
		1 (2.0)			Striatum
U92885	Efna3	apherin A3			Hypothalamus
			growth factor	Coton	Cortex
	1	1		Breast	Cortex
MM_013852	Cas	chemokine (C-C motif) ligand 4		Lung	Striatum
	1		growth factor	Colon	Striatum
	1			Breast	Cortex
	1	1			Hypothalamus
	1		i	Lung	Cortex
NM_009757	Bmp15	Growth differentiation factor the and the base	ŀ		Striatum
-		growth differentiation factor-9b gdf-9b; bone morphogenetic protein 15 bir	growth factor	Colon	Midbrain
	ì				Cortex
	1	1		Breast	Midbrain
	1				Cortex
	1	<u> </u>			Hypothalamus
	1		1	Lung	Midbrain
	1				Striatum
NM_008675	Nbi1	neuroblastoma, suppression of tumorigenicity t		i	Hypothalamus
-		the second of th	tumor related (secreted)	Colon	Midbrain
	1		1	1	Hypothatamus
	1		l	Breast	Cortex
			1	Lung	Midbrain
	1		J	Ì	Striatum
AK017955	Myg1	melanocyte prolifeating gene 1			H) pothalamus
	1	The style premoter of pene 1	signaling (secreted)	Colon	Midbrain
	1	i	l	( )	Cortex
	1	1	1	1	Striatum
	1		ļ		Hypothalamus
	ł		1	Breast	Hypothalamus
	1			Lung	Stratum
NK008922	Faf22	fibroblast growth factor 22			Hypothalamus
	1		growth factor	Colon	Striatum
	1			Breast	Cortex
	I				Hypothalamus
	1		Ì	Lung	Midbrain
	<del></del>		<u></u>		Cortex



#### **ABSTRACT**

The invention features methods and compositions for diagnosing non-central nervous system (non-CNS) disorders by detecting changes in gene expression in the CNS.

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